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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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16-AUG-1996;
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                                                                                                                                                                                                                                                    Compsn. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
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Matches 8; Conserv
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The invention relates to conjugate peptides engineered to noncovalently behind to heat shock proteins (hsp). A method of identifying a hsp binding peptide comprises (a) contacting a phage display library having bacteriophage expressing, in a surface protein, inserted peptides with a hsp target, and bound to a benzoquinone ansamycin antibictic (BAA), in a physiologic binding buffer; (b) isolating a phage binding to the hsp target; and (c) identifying the inserted peptide expressed. The peptides which bind to a hsp can be used as tethering peptides for a hsp which may serve as an accessory in a chaperone process and/or may comprise a cytokine. They can also be coupled to antigens to induce an immune response. Such compositions can be used for treating neoplastic disease, e.g. cancers, infectious diseases, e.g. diseases caused by a bacterium, virus, protozoan, mycoplasma, fungus, yeast, parasite or prion, or a missimmune system, e.g. acquired immune deficiencies or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a hear shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Melanoma antigen; MART-1; MAGE-1; gpl00; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; abGE; NYSSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 2e+06; cive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein and a melanoma antigen i.e. tyrosinase,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Houghton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Melanoma antigen, javelin peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAU72015 standard, peptide; 8 AA.
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(first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                                                                                                                                                           autoimmune diseases
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Best Local Similarity
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26-FEB-2002
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antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention. (Updated on 11-SEP-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunogenic complex; non-pathogenic multi-component viral particle; javelin; heat shock protein; humoral immunity; cellular immunity; anti-viral immune response; viral infection; hepatitis; influenza; mumps; HIV infection; human immunodeficiency virus; polio; tick-borne encephalitis; ebola virus infection; heat shock binding peptide.
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                                                                                                                                                                                              / Match 100.0%; Score 64; DB 4; Length 8; Local Similarity 100.0%; Pred. No. 2e+06; nee 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heat shock binding peptide #1 useful as javelin peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                 AAU80565 standard; peptide; 8 AA.
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                                                                                                                                                              Sequence 8 AA;
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Heat shock protein; hap; conjugate peptide; benzoquinone ansamycin; infectious disease; human papillomavirus; herpes virus; retrovirus; hepatitis virus; influenza virus; rhinovirus;

Heat shock protein (hsp) conjugate peptide #71

(first entry)

11-MAR-2004

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                                                                                                                                                                                                                                                                                                                                                                      A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                     Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy.
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100.0%; Score 64; DB 5; Length 8; 100.0%; Pred. No. 2e+06; ive 0; Mismatches 0; Indels
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                                                                                                 AAE13434 standard; peptide; 8 AA.
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                  Conservative
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MAYHEW M.
HOE M.
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Best Local Similarity
8; Conserve
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Matches 8; Conserv
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                                                                                                                  AAE13434;
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(HOEM/)
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The invention relates to a method for identification of a peptide binding to a heat shock protein (hsp) involving contacting a phage display contacted peptides with an hsp target or hsp target bound to a benzoquinone ansamycin antiblotic in a physiological binding buffer or binding buffer, isolating the phage, and identifying the inserted peptide expressed in the surface protein of the phage. The method is useful for identifying a peptide for inducing an immune response, for the treatment of infectious diseases (e.g. diseases caused by a bacterium, witcus, or infectious diseases (e.g. diseases acused by a bacterium, virus, or infectious diseases (e.g. diseases sursed by a bacterium, virus, or papillomavirus, a herpes virus such as herpes simplex virus, a retrovirus cuch as human immunodeficiency virus in 2, a hepatitis virus, a retrovirus cuch as human immunodeficiency virus, repiratory syncytial virus, a cytomegalovirus, an adenovirus, repiratory syncytial virus, a cytomegalovirus, an adenovirus, streptococcus, Enterococcus, Clostridium, Escherichia, Klebsiella, Vibrio or Mycobacterium, a protozoan such as an amoeba, malarial parasite or Trypanosoma cruzi), or carcinoma of the prostate, ovarian carcinoma, carcinoma of the prostate, ovarian carcinoma, carcinoma of the prostate, ovarian carcinoma of the breast.

Cu terine carcinoma, colon carcinoma, carcinoma of the breast.

Cu terine carcinoma, colon carcinoma, carcinoma of the lung, gilloblastoma or astrocytoma) or immunological diseases (e.g. AIDS) and for treating carcinoma carcinoma, carcinoma carcinoma, carcinoma or the prostate, ovarian carcinoma of the prostation and manamental diseases such as theumatodid arthritis, systemic lung.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identification of a peptide binding to a heat shock protein involves contacting a phage display library of several bacteriophages expressing in a surface protein of inserted peptides with a target followed by isolation and identification.
respiratory syncytial virus; cytomegalovirus; adenovirus; malignant disease; neoplastic disease; sarcoma; lymphoma; leukaemia; melanoma; carcinoma; breast; prostate; ovary; cervix; uterus; colon; lung; glioblastoma; astrocytoma; immunological disease; AIDS; autoimmune disease; rheumatoid arthritis; systemic lupus erythematosus; diabetes mellitus; thyroiditis; multiple sclerosis.
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OUERPELLI O.
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                                                                                                                                                                                                                                                                                                                  Synthetic.
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HWDFAWPW 8

ADG72529 standard; peptide; 8 AA.

RESULT 6 ADG72529 ADG72529

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                                                                                                                                                                                                                                                                                                                                                               infectious disease; human papillomavirus; herpes virus; retrovirus; hepatitis virus; influenza virus; rhinovirus; respiratory syncytial virus; cytomegalovirus; adenovirus; malignant disease; neoplastic disease; sarcoma; lymphoma; leukaemia; melanoma; carcinoma; breast; prostate; ovary; cervix; uterus; colon; lung; glioblastoma; astrocytoma; immunological disease; AIDS; autoimmune disease; rheumatoid arthritis; systemic lupus erythematosus; diabetes mellitus; thyroiditis; multiple sclerosis.
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              ; Score 64; DB 7; Length 8;
; Pred. No. 2e+06;
0; Mismatches 0; Indels
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                                                                                                                                                                                                          ADG72870 standard; peptide; 8 AA.
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                  100.0%;
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Best Local Similarity 100.
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HARTL U.
OUERFELLI O.
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influenza virus, a rhinovirus, respiratory syncytial virus, a cytomegalovirus, an adenovirus, Mycoplasma pneumoniae, a bacterium of the genus Salmonella. Staphylococcus, Streptococcus, Enterococcus, Clostridium, Escherichia, Klebalella, Vibrio or Mycobacterium, a protozoan such as an amoeba, malarial parasite or Trypanosoma cruzi), or malignant diseases, for treating or preventing neoplastic diseases (e.g. sarcoma, lymphoma, leukaemia, melanoma, carcinoma of the breast, carcinoma, of the prostate, ovarian carcinoma, carcinoma of the cervix, uterine carcinoma, colon carcinoma, carcinoma of the lung, glioblastom a satrofmenune diseases or immunological diseases (e.g. AIDS) and for treating autoimmune diseases uncolon arthritis, systemic lupus erythematosus, diabetes mellitus, thyroiditis and multiple sclerosis. This sequence represents an hsp related peptide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel hybrid antigen which comprises at leas one antigenic domain of an infectious agent or tumour antigen and a binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hybrid antigen; antigenic domain; infectious agent; tumour antigen;
binding domain; heat shock protein; antimicrobial; cytostatic; vaccine;
gene therapy; infectious disease; cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New hybrid antigens comprising an antigenic domain and improved heat shock protein-binding domains, useful for preventing or treating infectious diseases or cancer.
                                                                                                                                                                                                                                                                                                                                                       Gapa
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                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 7; Length 8; 100.0%; Pred. No. 2e+06; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel hybrid antigen-related peptide #1314.
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18-APR-2003; 2003US-0463746P.
16-SEP-2003; 2003US-0503417P.
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                                                                                                                                                                                                                                                                                                                                                     8; Conservative
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                                                                                                                                                                                                                                                                           Sequence 8 AA;
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The invention relates to hybrid antigens comprising at least one antigenic domain of an infectious agent or tumour antigen, at least one binding domain that non-covalently binds to a heat shock protein (hsp), and at least one peptide linker between them. Also disclosed are: (a) a composition for inducing an immune response to an infectious agent or tumour antigen comprising at least one of the hybrid antigens or a complex of at least one heat shock protein and at least one of the hybrid antigens, (b) a method for inducing an immune response to an infectious agent or tumour antigen by administering a hybrid antigen and a heat shock protein, where the hybrid antigen and the heat shock protein are non-covalently bound, and (c) treating an infectious disease or cancer by administering a hybrid antigen and a heat shock protein a preferably hsp70. The composition is administered via oral or protein is preferably hsp70. The composition is administered via oral or for treating or preventing cancer or infectious disease. The new peptide linkers give the antigens improved activity. Note: Many of the SEQ ID Nos
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infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; hap; immune response; infectious disease; cancer; cytostatic; antimicrobial; immunostimulant.
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                                                                                                                                                         Score 64; DB 8; Length 8;
Pred. No. 2e+06;
Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heat shock protein binding domain #226.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADU08477 standard; peptide; 8 AA.
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; 2003US-0503417P.
; 2004US-00776521.
; 2004WO-US004340.
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                                                                                                                                                                                                                                                                                             HWDFAWPW 8
                                                                                                                                                                                           Similarity
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                                                                                               Sequence 8 AA;
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16-SEP-2003;
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                                                                                                                                                         Query Match
Best Local S:
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid antigen useful for treating an infectious disease or cancer, comprises an antigenic domain from the infectious agent or cancer joined to a heat shock protein binding domain through an improved linker
are replicated more than once in the specification but the sequences of these replicated SEQ ID Nos are not the same. This sequence represents heat shock protein binding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; hsp; immune response; infectious disease; cancer; cytostatic; antimicrobial; immunostimulant.
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                                                                                                                  Length 8;
                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         High affinity heat shock protein binding sequence #1.
                                                                                                                  100.0%; Score 64; DB 8; 100.0%; Pred. No. 2e+06;
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                                                                                                                                                                                                                                                                                                                                  ADU07913 standard; peptide; 8 AA
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18-APR-2003; 2003US-0463746P.
16-SBR-2003; 2003US-0503417P.
13-FEB-2004; 2004US-00776521.
13-FEB-2004; 2004US-00820067.
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Query Match

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                                                                             Sequence 8 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified.
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Matches
                                                                              RESULT 12
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linkers give the antigens improved activity. Note: Many of the SEQ ID Nos are replicated more than once in the specification but the sequences of these replicated SEQ ID Nos are not the same. This sequence represents a heat shock protein binding sequence.
                                                                                                                                                                                                                                                                                                                                   Immunogenic complex; non-pathogenic multi-component viral particle; javelin; heat shock protein; humoral immunity; cellular immunity; anti-viral immune response; viral infection; hepatitis; influenza; mumps; HIV infection; human immunodeficiency virus; polio; tick-borne encephalitis; ebola virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New heat shock protein-based viral vaccines, useful for enhancing antiviral immune response in an organism, particularly as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza or HIV infection.
                                                                                                                 Gaps
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                                                                                       Query Match 100.0%; Score 64; DB 8; Length 8; Best Local Similarity 100.0%; Pred. No. 2e+06; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                              Javelin peptide #5 comprising javelin and linker regions.
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                                                                                                                                                                                                                                   AAU80739 standard; peptide; 9 AA.
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                                                                Sequence 8 AA;
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100.0%; Score 64; DB 5; Length 9; 100.0%; Pred. No. 2e+06; ive 0; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                  Immunogenic complex; non-pathogenic multi-component viral particle; javelin; heat shock protein; humoral immunity; cellular immunity; anti-viral immune respones; viral infection; hepatitis; influenza; mumps; HIV infection; human immunodeficiency virus; polio; tick-borne encephalitis; ebola virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New heat shock protein-based viral vaccines, useful for enhancing antiviral immune response in an organism, particularly as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza or HIV infection.
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                                                                                                                                                                                                                                                                                                                                                            Javelin peptide #6 comprising javelin and linker regions.
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tick-borne encephalitis; ebola virus infection
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Synthetic.
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                                                                          25-OCT-2001
                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to the use of an immunogenic complex, comprising a non-pathogenic multi-component viral particle covalently linked to a javalin molecule (preferably a peptide) that selectively binds to a heat shock protein. The immunogenic complex is useful for inducing both humoral and cellular immunicy, especially for enhancing the anti-viral immune response, in a human or non-human subject. The immunogenic complex is particularly useful as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza, mumps, HIV or ebola virus infection, anusuodeficiency virus) infection, polio, tick-borne encephalitis or ebola virus infection. ANUS0735-AAUS0742 represent javelin neptides, comprising javelin and linker regions, which may be covalently conjugated to multi-component viral particles
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                                                                      Immunogenic complex; non-pathogenic multi-component viral particle; javelin; heat shock protein; humoral immunity; cellular immunity; arti-viral immune response; viral infection; hepatitis; influenza; mumps; HIV infection; human immunodeficiency virus; polio; tick-borne encephalitis; ebola virus infection.
                                                                                                                                                                                                                                                                                                                                                   New heat shock protein-based viral vaccines, useful for enhancing antiviral immune response in an organism, particularly as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunogenic complex; non-pathogenic multi-component viral particle; javelin; heat shock protein; humoral immunity; cellular immunity; anti-viral immune response; viral infection; hepatitis; influenza; mumps;
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                                                 Javelin peptide #1 comprising javelin and linker regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Javelin peptide #2 comprising javelin and linker regions.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 14; 75pp; English.
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                                                                                                                                                                                                                                                                            (MOJA-) MOJAVE THERAPEUTICS INC.
                                                                                                                                                                                                                            17-APR-2001; 2001WO-US012568,
                                                                                                                                                                                                                                                     17-APR-2000; 2000US-0197462P
                        (first entry)
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|HWDFAWPW 12
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                                                                                                                                                                                                                                                                                                                                                                                           HIV infection.
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AAU80735
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comprising a non-pathogenic multi-component viral particle covalently linked to a javelin molecule (preferably a peptide) that selectively binds to a heat shock protein. The immunogenic complex is useful for inducing both humoral and cellular immunity, especially for enhancing the anti-viral immunosenic complex is particularly useful as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza, mumps, HIV (human immunodeficiency virus) infection, polio, tick-borne encephalitis comprising javelin and linker regions, which may be covalently conjugated to multi-component viral particles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New heat shock protein-based viral vaccines, useful for enhancing antiviral immune response in an organism, particularly as a vaccine for preventing or ameliorating viral infections, e.g. hepatitis, influenza or HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 14; 75pp; English.
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                                                                                                                                                                                                                                                                                                (MOJA-) MOJAVE THERAPEUTICS INC.
                                                                                                                                                                                                                      17-APR-2000; 2000US-0197462P.
                                                                                                                                                17-APR-2001; 2001WO-US012568.
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WO200178772-A1
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigens, GMZ, antigens, GMZ, antigens, GMZ, antigens of the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANJ1980-AAU72481 represent melanoma antigen peptides of the invention
                                            cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                           100.0%; Score 64; DB 4; Length 16; 100.0%; Pred. No. 0.0068; ative 0; Mismatches 0; Indel8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti cancer vaccine for the treatment of melano protein and a melanoma antigen i.e. tyrosinase.
                                                            protein and a melanoma antigen i.e. tyrosinase
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                                                                                          Disclosure, Page 25, 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU72188 standard; peptide; 17 AA.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                               1 HWDFAWPW 8
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             WPI; 2001-663092/76
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                                                                                                                                                                                                                                                                                                                 Sequence 16 AA;
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(ALAW/) AL-AWQA
(MAYH/) MAYHEW
(HOEM/) HOE M.
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                                                                                                                                                                                    cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                           Mayhew M,
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                                                                                                                                                                                                      protein and a melanoma antigen i.e. tyrosinase
                                                                                                                         Al-Awgati Q,
                                                                                                                                                                                                                                     Disclosure; Page 25; 150pp; English
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17-APR-2000; 2000US-0197462P
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                                                                                                                         Livingston P,
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAXH/) MAXHEW M.
                           HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                           HOE M.
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antigen peptides of the invention
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           a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp.100, MAGB antigens, BAGB antigens, NYESOL, MART antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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comprising administration of an immunotherapeutic composition, comprising
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; WSES01; GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                                                                                                                                                            100.0%; Score 64; DB 4; Length 17; 100.0%; Pred. No. 0.0072;
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                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                             AAU72227 standard; peptide; 17 AA.
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LIVINGSTON P.
AL-AWQATI Q.
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Best Local Similarity
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100.0%; Score 64; DB 4; Length 17; 100.0%; Pred. No. 0.0072; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                             AAU72296 standard; peptide; 17 AA.
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                                                               8; Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Query Match
Best Local Similarity
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigens, GMZ, antigens, OFTLOS antigens and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AdVI2481 represent melanoma antigen peptides of the invention
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                                                                                                        Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immuncherapeutic; heat shock protein; tyrosinase hagE; WRSB01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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100.0%; Pred. No. 0.0072;
tive 0; Mismatches 0; Indels
                                                                 MAGE-3-derived melanoma antigen, javelin peptide #6.
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                      26-FEB-2002 (first entry)
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(LIVI/) LIVINGSTON P
(ALAW/) AL-AWQATI Q.
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(ALAW/) AL-AWQI
(MAYH/) MAYHEW
(HOEM/) HOE M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
HOE M.
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(ALAW/) (MAYH/) (HOEM/)

(HOUG/) (LIVI/)

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26-FEB-2002 (first entry)
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                              HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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(LIVI/) LIVINGSTON P.
(ALAW) AL-AWQATI Q.
(MAXH/) MAXHEW M.
(HOEM/) HOE M.
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(MAYH/)
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                                                                                 17-APR-2001; 2001WO-US012449.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                                                                    (HOEM/) HOE M.
            sapiens.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a heat shock
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                                                                                                                                                                                                                                                                                                                                                      Houghton A, Livingston P, Al-Awgati Q,
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17-APR-2000; 2000US-0197462P.
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invention relates to a method of induction of an immune response.
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                                                                                                        comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, WART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                             treatment of melanoma comprises a heat shock
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    Mayhew M,
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                                                        a melanoma antigen i.e. tyrosinase
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    Al-Awgati Q,
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                                                                          Disclosure; Page 31; 150pp; English
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                                                                                                                                                                                                       antigen peptides of the invention
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                                         Anti cancer vaccine for the
protein and a melanoma antio
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    Livingston P,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                        WPI; 2001-663092/76
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Best Local Similarity
Matches 8; Conserv
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    Houghton A,
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NVES01, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the bat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAU72481 represent melanoma antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2.
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
Javelin molecule; melanoma antigen recognised by T cells-1; human.
for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.0072;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                          4; Length 17;
                                                        Query Match
100.0%; Score 64; DB 4; Length 17
Best Local Similarity 100.0%; Pred. No. 0.0072;
Matches 8; Conservative 0; Mismatches 0; Indels
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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HWDFAWPW 17
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                                                                                                                                                                                                                                                                                                   Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; ABGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                MAGE-1/3-derived melanoma antigen, javelin peptide #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Houghton A, Livingston P, Al-Awqati Q, Mayhew M,
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AAU72447 standard; peptide; 17 AA.
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                                                                                                                                                     (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, ratigence, BAGE antigens, NYESQ1, MART antigens, GM2, antigence portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS481 represent melanoma antigen peptides of the invention
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javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                            Al-Awqati Q,
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                                                                                                                                             17-APR-2001; 2001WO-US012449.
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                                                                                                                                                                                                                                                                                                            Livingston P,
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                                                                                                                                                                                                            (HOUG/) HOUGHTON A.
(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Best Local Similarity
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                                                                               WO200178655-A2
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                              Homo sapiens.
Synthetic.
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(HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                            Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                Tyrosine-derived melanoma antigen, javelin peptide #1.
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                                26-FEB-2002 (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                               Homo sapiens.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule; ann-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANT1980-ANU72481 represent melanoma antigen peptides of the invention
                                                                                   melanoma comprises a heat shock
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                                                                                                                                                                       The invention relates to a method of induction of an immune response.
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                  Mayhew M,
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                                                                                  Anti cancer vaccine for the treatment of melano protein and a melanoma antigen i.e. tyrosinase.
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                  Al-Awgati Q,
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                  Livingston P,
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Best Local Similarity 100.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Synthetic.
                  Houghton A,
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(MAYH/)
(HOEM/)
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                                                                                                                                                                                                                                                           protein and a melanoma antigen i.e. tyrosinase
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17-APR-2001; 2001WO-US012449
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Matches 8; Conservative
                                                                 HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                (LIVI/) LIVINGS
(ALAW/) AL-AWQA
(MAYH/) MAYHEW
(HOEM/) HOE M.
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Synthetic.
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(MAYH/)
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Gaps

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Melanoma antigen, MART-1, MAGE-1, gp100, cytostatic; immune response; immuncherapetic; heat shock protein; tyrosinase; BAGE; NYSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; favelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU72277 standard; peptide; 17 AA.
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LIVINGSTON P.
AL-AWQATI Q.
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shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                       100.0%; Score 64; DB 4; Length 17; 100.0%; Pred. No. 0.0072; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                  AAU72406 standard; peptide; 17 AA.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI O.
(MAYH/) MAYHEW M.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                    MART-1-derived melanoma antigen, javelin peptide #6.
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Best Local Similarity 100.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72396
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AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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HOE M.

(ALAW/) (MAYH/) (HOEM/)

(HOUG/)

Houghton A,

RESULT 37

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Homo sapiens

Synthetic

25-OCT-2001

26-FEB-2002

AAU72476;

RESULT 36

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Hoe M;

(HOUG/)

(HOEM/) (ALAW/) (MAYH/)

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                     17-APR-2001; 2001WO-US012449.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                       HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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                                                                                                     (HOUG/)
(LIVI/)
(ALAW/)
(MAYH/)
(HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEGO1, MART antigen is covalently bound and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTISBO-AAUTISBI represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                         doughton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 30; 150pp; English
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                   Homo sapiens
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                                                                                      Synthetic.
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Homo sapiens.
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                                                                                                                       comparising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AMV71980-AAMV72481 represent melanoma antigen peptides of the invention
                                                            comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the treatment of melanoma comprises a heat shock
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                                                                                                              The invention relates to a method of induction of an immune response
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                     Hoe M;
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                                                                                                                                                                                                                                                               100.0%; Score 64; DB 4; Length 17; 100.0%; Pred. No. 0.0072;
                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            lyrosine-derived melanoma antigen, javelin peptide #20.
                     Mayhew M,
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                                                             melanoma
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protein and a melanoma antigen i.e. tyrosinase.
                                                            Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase.
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                    Livingston P, Al-Awgati Q,
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                                                                                           Disclosure; Page 33; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                AAU72206 standard; peptide; 17 AA.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                       WPI; 2001-663092/76
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(HOEM/) HOE
                     Houghton A,
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                                                          The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antiegen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat for the treatment of melanoma antigen peptides of the invention AMJ1980-AMJ7481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 4; Length 17; 100.0%; Pred. No. 0.0072;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tyrosine-derived melanoma antigen, javelin peptide #31.
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Disclosure; Page 23; 150pp; English.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 AA;
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherspeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEsO1, MART antigen for antigen for these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANTIJS80-AAUT1281 represent melanoma antigen peptides of the invention
antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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Best Local Similarity 100.
Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                          Sequence 17 AA;
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(HOEM/)
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(ALAW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprises a heat shock
                                                                                                                                                                                                                                     Melanoma antigen; MART-1; MAGB-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
Gapa
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Indele
                                                                                                                                                                                                              MART-1-derived melanoma antigen, javelin peptide #36
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Mismatches
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Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Synthetic.
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(MAYH/) MAYHE
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RESULT 45 AAU72236

Score 64; DB 4; Length 17; Pred. No. 0.0072;

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Query Match Best Local Similarity

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HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAXHEW M.
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                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                               Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase abdGE, WYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                            gp100-derived melanoma antigen, javelin peptide #15.
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                                                                                                                                                                                                                                                                                                                                                                protein and a melanoma antigen i.e. tyrosinase
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AAU72236 standard; peptide; 17
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigen protein 2, gp 100, MAGE antigens, BAGE antigens of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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    BAGE; NYES01; GM2;
immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2 tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                   Houghton A, Livingston P,
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(MAYH/) MAYHEW M.
(HOEM/) HOE M.
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                                                                                                                                                                            Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                    Mayhew M,
                                                                                                                                   Livingston P, Al-Awqati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
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Synthetic.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                 melanoma comprises a heat shock
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Best Local Similarity 100.0%; Pred. No. 0.0072;
Matches 8; Conservative 0; Mismatches 0; Indels
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маурем м,
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protein and a melanoma antigen 1.e. tyrosinase
Al-Awgati Q,
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antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention

Sequence 18 AA;

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melanoma

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase ratigens, BAGE antigens, NYESO1, WART antigens, CM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the bat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUT1980-AAUT2481 represent melanoma antigen peptides of the invention
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                                                method of induction of an immune
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                 Disclosure; Page 28; 150pp; English.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                The invention
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 100.0%; Score 64; DB 4; Length 18; 100.0%; Pred. No. 0.0076;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer vaccine for the treatment of
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                                                                                                                                                                   AAU72326 standard; peptide; 18 AA.
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              Local Similarity 100.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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MAGE-1-derived melanoma antigen, javelin peptide #26.
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(LIVI/) LIVINGSTON P.
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LIVINGSTON P.
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MAYHEW M.
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(ALAW/)
(MAYH/)
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AAU72456
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                               Homo sapiens
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(HOEM/)
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Best Loca Matches

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                                                                                                                             The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigent c portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANU71980-AAU712481 represent melanoma antigen peptides of the invention
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                                                                                 comprises a heat shock
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                                        Mayhew M,
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                                                                                 melanoma
                                                                               Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase
                                        Al-Awgati Q,
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                                                                                                              Disclosure; Page 33; 150pp; English
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Matches 8; Conservative
                                        Livingston
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
AL-AWQATI Q.
MAYHEW M.
HOE M.
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                                                            WPI; 2001-663092/76
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                                        Houghton A,
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(HOEM/)
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(HOEM/)
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                                                                                                               The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, ratigens, BAGE antigens, NYED1, MART antigens, GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 64; DB 4; Length 18; 100.0%; Pred. No. 0.0076; ative 0; Mismatches 0; Indels
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protein and a melanoma antigen i.e. tyrosinase
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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nes 8, Conserv
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Best Local Similarity Matches 8; Conser

Query Match

AAU72457 standard; peptide; 18 AA

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AAU72457;

100.0%; Score 64; DB 4; Length 18; 100.0%; Pred. No. 0.0076;

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              antigen is covalently bound to a javelin molecule, where the melanoma stringen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                                                                                                                          100.0%; Score 64; DB 4; L 100.0%; Pred. No. 0.0076; Live 0; Mismatches 0;
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 GM2, antigenic portions
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                                                                            antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                             Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    melanoma comprises a heat shock
                                                                                                                                   Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapetic; heat shock protein; tyrosinase; BAGE; NYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; tyrosinase related protein 2; vaccine; tyrosinase protein 2; vaccine; tyrosinase related protein 1; vaccine; tyrosinase related by T cells-1; human.
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                                                                                                 MAGE-1/3-derived melanoma antigen, javelin peptide #16.
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                                                               (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Sequence 18 AA;

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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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                                                                                                                                                                                                                                                                                                                                                                                                 Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                        Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                     gp100-derived melanoma antigen, javelin peptide #25.
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                     AAU72246 standard; peptide; 18 AA
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
Matches 8; Conserv
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Synthetic.
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RESULT 61
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the bat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; NYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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Hoe M;

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                                                                 Al-Awgati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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WO200178655-A2
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                                                                              The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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cancer vaccine for the treatment of melanoma comprises a heat shock
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                protein and a melanoma antigen i.e. tyrosinase.
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                                                  Disclosure, Page 33; 150pp; English
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the pavelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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(LIVI/) LIVINGSTON P
(ALAW/) AL-AWQATI Q.
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSB01; GN2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                      Jp100-derived melanoma antigen, javelin peptide #135.
                                                 AAU72356 standard; peptide; 18
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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17-APR-2001; 2001WO-US012449.
                                                     17-APR-2000; 2000US-0197462P.
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                                                                                                        HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
                                                                                                                                                                                                                                                                                                                           NPI; 2001-663092/76
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                                                                                                                                                                                         MAYHEW M.
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1 HWDFAWPW 8

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HWDFAWPW

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Sequence 18 AA;
                   WO200178655-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                               Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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MAGE-1-derived melanoma antigen, javelin peptide #36.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU72287 standard; peptide; 18 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                          HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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11 HWDFAWPW 18
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Best Local Similarity
Matches 8; Conserv
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Synthetic.
                                                                                                                      Homo sapiens
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                                                                                                                                    Synthetic.
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(MAYH/)
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NESG1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                                                       Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer vaccine for the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Page 26; 150pp; English
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17-APR-2001; 2001WO-US012449.
                                                                     17-APR-2000; 2000US-0197462P.
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ses 8; Conservative
                                                                                                                                           (HOUG/) HOUGHTON A.
(LIVI/) LIVINGSTON P.
(ALAW) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                                                  Sequence 18 AA;
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Matches
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                                                                                                                                                 The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GM2, antigence for the selected protein 2, gp 100, MAGE antigens, BAGE antigens, NYEsO1, MART antigen is covalently bound on a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat for the treatment of melanoma is useful for inducing an immune response for the treatment of melanoma AMVI1980-AAVI2481 represent melanoma antigen peptides of the invention
                                                                                                  Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                            Hoe
                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gp100-derived melanoma antigen, javelin peptide #126.
                                                            Mayhew M,
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                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                            Al-Awgati Q,
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                                                                                                                               Disclosure; Page 27; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                         AAU72347 standard; peptide; 18 AA
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                                                            foughton A, Livingston P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                               WPI; 2001-663092/76
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                                        HOE M
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(ALAW/)
(MAYH/)
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(ALAW/)
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Anti cancer vaccine for the treatment of melanoma comprises a heat shock
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immunotherapeutic; heat shock protein; tyrosinase; aGGE; WSES01; GM2;
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                          protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                       Disclosure; Page 28; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-2002 (first entry)
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Les 8; Conservative
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Gape

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Indele

Pred. No. 0.0081; Mismatches 0;

100.0%; Pre-

Conservative

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12 HWDFAWPW 19
                               1 HWDFAWPW 8
  Best Local Similarity
                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                         Peptide
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Matches
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is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEsOl, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUT1980-AAUT2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compsn. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid peptides OVA-BiP (AAW19956) and BiP-OVA (AAW19957) comprise chicken OVA-peptide (see AAW19955) joined via a peptide linker to heat benck protein (HSP) BiP binding domain (see also AAW19951). The hybrid peptide can be combined in vitro with a HSP, such as hsp70, to form a complex that, when administered to a subject, induces an immune response
                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                             immunotherapy; heat shock protein; BiP; OVA; cancer;
                                                                                                               100.0%; Score 64; DB 4; Length 18;
100.0%; Pred. No. 0.0076;
ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Takeuchi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Houghton A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                                                                                                                                                                                                                          OVA-BiP-binding domain hybrid peptide.
                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                             AAW19956 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 18; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hartl FU, Hoe MH,
                                                                                                                                                                                                                                                                                                                                                                                               9. .11
/label= Linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-0002479P.
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                                                                                                                                                                                                                                                                                                                                                                           1. .8
/label= OVA
                                                                                                                                                                                                                                                                                                                                                                                                                   12. .19
/label= BiP
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                   Conservative
                                                                                                                                                                            11 HWDFAWPW 18
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                                                                                                                      Local Similarity
wes 8; Conserv
                                                                                                                                                                                                                                                                                                                         infectious disease.
                                                                                                                                                      1 HWDFAWPW
                                                                                             Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-AUG-1995;
18-AUG-1995;
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                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                  AAW19956;
                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                         Peptide
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Matches
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Length 19

100.0%; Score 64; DB 2;

Sequence 19 AA;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid peptides OVA-Bip (AAW19956) and Bip-OVA (AAW19957) comprise chicken OVA-peptide (see AAW19955) joined via a peptide linker to heat shock procein (HSP) Bip binding domain (see also AAW19951). The hybrid peptide can be combined in vitro with a HSP, such as hsp70, to form a complex that, when administered to a subject, induces an immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                      Vaccine; immunotherapy; heat shock protein; B1P; OVA; cancer; infectious disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 64; DB 2; Length 19; 100.0%; Pred. No. 0.0081; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SLOK ) SLOAN KETTERING INST CANCER RES
                                                                                                                              BiP-binding domain-OVA hybrid peptide
                                                                                                                                                                                                                                                                                    Location/Qualifiers
AAW19957 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU72254 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 18; 58pp; English.
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/label= Linker
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                                                                                                                                                                                                                                                                                                              . .8
label= BiP
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/label= Ova
                                                                                     10-NOV-1997 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-165035/15.
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18-AUG-1995;
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                                                                                                                                                                                                                                             Synthetic.
                                    AAW19957;
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                                                                                                                                                                                                                                                                                           Key
Peptide
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ID AAU7
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us-10-053-520-143.rag

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javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                      17-APR-2001; 2001WO-US012449.
                                                                                                                           L7-APR-2000; 2000US-0197462P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; Conservative
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/note=
                                                                                                                                                 HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                                                                                                                                                            IPI; 2001-663092/76
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 AA;
                                                         40200178655-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gallus gallus
                         Homo sapiens
                                                                                35-OCT-2001
                                    Synthetic.
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                                                                                                                                                 (HOUG/)
                                                                                                                                                                                                (HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, pp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
                                                                Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                           Hoe M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 19; 100.0%; Pred. No. 0.0081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                             JP100-derived melanoma antigen, javelin peptide #33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gp100-derived melanoma antigen, javelin peptide #32
                                                                                                                                                                                                                                                                                                                           Houghton A, Livingston P, Al-Awgati Q, Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 25; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU72253 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antigen peptides of the invention
                                                                                                                                                                                                        17-APR-2001; 2001WO-US012449.
                                                                                                                                                                                                                                17-APR-2000; 2000US-0197462P
                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                     HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HWDFAWPW 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 HWDFAWPW 8
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les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 AA;
                                                                                                                                                            WO200178655-A2
                                                                                                                                                                                                                                                                                                     HOE M.
                                                                                                                           sapiens
                      26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2002
                                                                                                                                                                                   25-OCT-2001
                                                                                                                                       Synthetic.
AAU72254;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                       (HOUG/)
                                                                                                                                                                                                                                                                             (ALAW/)
(MAYH/)
                                                                                                                                                                                                                                                                                                     (HOEM/)
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Matches
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The invention relates to a method of induction of an immune response, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, related protein 1, tyrosinase related protein 2, 9p 100, MAGE antigens, BAGE antigens, NYE801, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javalin molecule, where the melanoma antigen bound to the javalin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                 Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy; chicken.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Hoe
Mayhew M,
                                                                                                                                                                                                                   protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "MHC class I epitope"
doughton A, Livingston P, Al-Awqati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chicken MHC class I peptide antigen #1.
                                                                                                                                                                                                                                                                                                   Disclosure; Page 25; 150pp; English
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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is Herpes simplex virus MHC class I peptide antigen
                                                                                                                                                                                                                                                                                                                                                             A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy; chicken.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chicken MHC class I peptide antigen #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure, Page 14, 47pp, English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAE13447 standard; peptide; 19 AA.
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                                                                                                                                                                                                                                                Hoe M;
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17-APR-2001; 2001WO-US012567.
                                                            17-APR-2000; 2000US-0197462P.
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                                                                                                                                                                                                                                                Rothman JE, Mayhew M,
                                                                                                                      (ROTH/) ROTHMAN J E. (MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                          WPI; 2002-017594/02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWDFAWPW 8
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Best Local Similarity
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                                                                                                                                                                                   HOEM/) HOE M
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   PAX PX X BX X PX F F F F F F F F X S X X W X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
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/note= "MHC class I epitope"
   /note= "Javelin sequence"
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/note= "Javelin sequence"
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/note= "Linker"
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                                                                                                                                                                                                                                                17-APR-2000; 2000US-0197462P
                                                                                                                                                                                      17-APR-2001; 2001WO-US012567
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Best Local Similarity 100.00
Berling 8; Conservative
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                                                                                                                                                                                                                                                                                                          (ROTH/) ROTHMAN J E. (MAYH/) MAYHEW M. (HOEM/) HOE M.
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                                                            WO200179259-A1
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Region Region Region

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A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                          Disclosure; Page 14; 47pp; English.
                                                                                                                                                                                                                                                                                                       ADG72871 standard; peptide; 19 AA.
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                                                                                                                                                                                                             8; Conservative
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                                                                                                                                                                                                                                                     12 HWDFAWPW 19
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HOUGHTON A.
HARTL U.
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                                                                                                                                                                   Sequence 19 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                            ADG72871;
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(MAYH/) N
(HOEM/) I
(HOUG/) I
(HART/) (OUER/)
(MORO/)
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                                                                                                                                                                                                                                                                                    RESULT 83
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                                                                                                               comprising a
                                                                                                                       The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken MHC class I peptide antigen
                                                            A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     tigenic complex; epitope; heat shock protein; HSP; tether; javelin; jor histocompatability complex; MHC; therapy; immune response;
                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                    100.0%; Score 64; DB 5; Length 19; 100.0%; Pred. No. 0.0081; tive 0; Mismatches 0; Indels
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/note= "MHC class I epitope"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                    Disclosure; Page 13; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                   AAE13453 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Linker"
                     Hoe M;
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                     Mayhew M,
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                                         WPI; 2002-017594/02
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antigenic complex;
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HOE M.
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                                                                                                                                                                                                                                 Sequence 19 AA;
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 (HOEM/) HOE
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                     Rothman JE,
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Region
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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is Herpes simplex virus MHC class I peptide antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Example 1; Page 38; 56pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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 Barber B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invertion Featers to a method for its paperide binding to a heat shock protein (hap) involving contacting a phage display to a library of several bacteriophage which express, in a surface protein, several inserted peptides which an hap target or hap target bound to a benzoquinone ansamycin antibiotic in a physiological binding buffer or binding buffer, isolating the phage, and identifying the inserted peptide expressed in the surface protein of the phage. The method is useful for identifying a peptide, which binds to a heat shock protein, which is used in a conjugate peptide for inducing an immune response, for the treatment of in a conjugate peptide for inducing an immune response, for the treatment of in a conjugate peptide for inducing an immune response, for the treatment of papillomavirus, a herpes virus such as herpes simplex virus, a retrovirus such as human immunodeficiency virus in 2, a hepatitis virus, an influenza virus, a rathnovirus, respiratory syncytial virus, an conformation and an adenovirus, respiratory syncytial virus, an influenza virus, a rathnovirus, streptococcus, Enterococcus, Covernidum, Bachericinia, Klabaiella, Vibrio on Mycobacterium of the protezon such as an amoeba, malarial parasite or Trypanosoma cruzi), or malignant diseases, for treating or preventing neoplastic diseases (correctionma of the prostate, ovarian carcinoma, carcinoma of the brease.

Correctionma of the prostate, ovarian carcinoma, carcinoma of the cervix, cuterine carcinoma, or immunological diseases (e.g. AlDS) and for treating autoimmune diseases such as rheumatoid arthritis, systemic lupus carcinom of expression carcinoma of the invention.

Correctionment of seases such as rheumatoid arthritis, systemic lupus carcinom of the sequence represents an hap related peptide of the invention.
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                                                        invention relates to a method for identification of a peptide binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; antimicrobial; cytostatic; vaccine; gene therapy; infectious disease; cancer.
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                             Example; Page 18; 62pp; English
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11.APR-2003; 2003US-0462469P.
18-APR-2003; 2003US-0463746P.
16-SEP-2003; 2003US-0503417P.
12-FEB-2004; 2004US-00776521.
isolation and identification.
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8; Conservative
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12 HWDFAWPW 19
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shock protein-binding domains, useful for preventing or treating
infectious diseases or cancer.
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18-APR-2003; 2003US-0463746P
16-SEP-2003; 2003US-0503417P
12-FEB-2004; 2004US-00776521
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B; Conservative
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WPI; 2004-625768/60
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                            one antigenic domain of an infectious agent or tumour antigen and a binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; antimicrobial; cytostatic; vaccine; gene therapy; infectious disease; cancer.
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This invention relates to a novel hybrid antigen which comprises
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Matches 8; Conserv
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16-SEP-2003;
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 Length 19;
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                         0; Indels
100.0%; Score 64; DB 8;
ilarity 100.0%; Pred. No. 0.0081;
Conservative 0; Mismatches 0;
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18-AUG-1995;
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                                                                                             Vaccine; immunotherapy; heat shock protein; BiP; human papilloma virus;
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                                                                            4PV hybrid protein BiP-E7 (Type 16) for use in vaccine.
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                                                                                                                                        Location/Qualifiers
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                       AAW19961 standard; peptide; 20 AA.
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HWDFAWPW
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                                                                                                                                                                                                                                                16-AUG-1996;
                                                                                                                                                                                                                                                                           18-AUG-1995;
                                                          10-NOV-1997
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                                                                                                                        Synthetic
                                        AAW19961;
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Peptide
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AAW19967
ID AAW199
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Compsn. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
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/accine; immunotherapy; heat shock protein; BiP; human papilloma virus;
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/label= Linker
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/label= BiP
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Compan. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
                                                                                                                                                                                                                                                    virus (HPV) comprise an HPV antigenic peptide joined via a peptide linker to a heat shock protein BiP binding domain (see also AAW19951). The peptide given in AAW19962 comprises a fusion of HCV E7 (Type 18) peptide and BiP binding domain. The hybrid peptide is combined in virro with a HSP to form a complex that, when administered to a subject, induces an
                                                                                                                                                                                                                                     Hybrid peptides (AAW19958-67) for use in vaccines against human papilloma
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                                                                               Houghton A,
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                                               (SLOK ) SLOAN KETTERING INST CANCER RES
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                                                                               Hoe MH,
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/label= Linker
13. .20
/label= BiP
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 95US-0002479P.
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/label= E7
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Best Local Similarity
8; Conserv?
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   18-AUG-1995;
                 18-AUG-1995;
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/label= Linker
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HWDFAWPW 20
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18-AUG-1995;
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Hybrid peptides (AAW19958-67) for use in vaccines against human papilloma virus (HPV) comprise an HPV antigenic peptide joined via a peptide linker to a heat shock protein BiP binding domain (see also AAW19951). The peptide given in AAW19959 comprises a fusion of BiP binding domain and HCV B7 (type 11) peptide. The hybrid peptide is combined in vitro with a HSP to form a complex that, when administered to a subject, induces an immune response against HCV
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a HSP to form a complex that, when administered to a subject, induces an immune response against \ensuremath{\mathsf{HCV}}
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100.0%; Pred. No. 0.0085;
iive 0; Mismatches 0; Indels
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                                                                                       100.0%; Score 64; DB 2;
100.0%; Pred. No. 0.0085;
iive 0; Mismatches 0;
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                                                                                     Hybrid peptides (AAW19958-67) for use in vaccines against human papilloma virus (HPV) comprise an HPV antigenic peptide joined via a peptide linker to a heat shock protein BiP binding domain (see also AAW19951). The peptide given in AAW19964 comprises a fusion of HCV E7.1 (Type 6b) peptide and BiP binding domain. The hybrid peptide is combined in vitro with a HSP to form a complex that, when administered to a subject, induces an immune response against HCV
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 protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
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; Pred. No. 0.0085;
0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                          Example 10; Page 24; 58pp; English.
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/label= Linker
12. .20
/label= E7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW19965 standard; peptide; 20
                                                                                                                                                                                                                                                                    100.0%;
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95US-0002490P
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/label= BiP
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Best Local Similarity 100.v.
Bertas 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                   13 HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                        1 HWDFAWPW 8
                                                                                                                                                                                                                                   Sequence 20 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW19965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
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AAW19963 ID AAW

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Hybrid peptides (AAW19958-67) for use in vaccines against human papilloma virus (HPV) comprise an HPV antigenic peptide joined via a peptide linker to a heat shock protein BiP binding domain (see also AAW19951). The peptide given in AAW19958 comprises a fusion of HCV E7 (type II) peptide and BiP binding domain. The hybrid peptide is combined in vitro with a HSP to form a complex that, when administered to a subject, induces an immune response against HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compsn. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment
                                                                      Vaccine; immunotherapy; heat shock protein; BiP; human papilloma virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYE801; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 100.0%; Score 64; DB 2; Length 20; Local Similarity 100.0%; Pred. No. 0.0085; e8 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Takeuchi Y,
                                    APV hybrid protein E7 (Type 11)-BiP for use in vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MAGE-1-derived melanoma antigen, javelin peptide #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hoe MH, Houghton A,
                                                                                                                                                                                                                                                                                                                                                                                                                                              (SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infectious diseases and tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Page 23; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU72404 standard; peptide; 20 AA
                                                                                                                                                                                                                        Linker
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95US-0002490P
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/label= BiP
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/label= E7
   10-NOV-1997 (first entry)
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/label= 1
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                                                                                                                       Synthetic
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                                                                                                                                                                                                        Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus (HPV) comprise an HPV antigenic peptide joined via a peptide linker to a heat shock protein BiP binding domain (see also AAW19951). The peptide given in AAW19963 comprises a fusion of BiP binding domain and HCV B7 (type 18) peptide. The hybrid peptide is combined in vitro with a HSP to form a complex that, when administered to a subject, induces an immune response against HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compsn. for inducing immune response contg. antigen and heat shock protein - also new hybrid peptide and related nucleic acid, for treatment of infectious diseases and tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid peptides (AAW19958-67) for use in vaccines against human papilloma
                                                                                                                                                                                                                                        Jaccine; immunotherapy; heat shock protein; BiP; human papilloma virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 100.0%; Score 64; DB 2; Length 20; Similarity 100.0%; Pred. No. 0.0085; 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                        HPV hybrid protein BiP-E7 (Type 18) for use in vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Houghton A,
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                                                                                                       AAW19963 standard; peptide; 20
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/label= Linker
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/label= BiP
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/label= E7
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HWDFAWPW
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SLOK ) SLOAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MO9706821-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-AUG-1996;
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                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                     AAW19963;
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Peptide
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Mayhew M;

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Gaps

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RESULT 96

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AAW19958 ID AAW1 XX AC AAW1 XX

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Houghton A, Livingston
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
         HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                13 HWDFAWPW 20
                                                                                            WPI; 2001-663092/76
                                                                                                                                                                                                                                                                                                                                                               1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                               Sest Local Similarity
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                                                                                                                                                                                                                                                                                                                       Query Match
                     (LIVI/)
(ALAW/)
(MAYH/)
(HOEM/)
           HOUG/)
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                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                               The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GA2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Melanoma antigen, MART-1, MAGB-1; gp100; cytostatic; immune response; immuncherapetic; heat shock protein; tyrosinase; BAGB; NYSSO1, GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                       Houghton A, Livingston P, Al-Awgati Q, Mayhew M, Hoe M;
                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; Arive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MAGE-1/3-derived melanoma antigen, javelin peptide #3.
                                                                                                                                                                                                                         protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                             Disclosure; Page 31; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                          antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU72444 standard; peptide; 20
                                                             17-APR-2001; 2001WO-US012449.
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                                                                                  17-APR-2000; 2000US-0197462P
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                                                                                                                                                                                                                                                                                                                                                                                                                                        8; Conservative
                                                                                                      HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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13 HWDFAWPW 20
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HWDFAWPW
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 AA;
                     WO200178655-A2
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                                          25-OCT-2001
 Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU72444;
                                                                                                                           (ALAW/)
                                                                                                                                                (HOEM/)
                                                                                                      (HOUG/)
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                                                                                                                                        melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Melanoma antigen; MART-1; MAGE-1; gpl00; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                               The invention relates to a method of induction of an immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085;
Hoe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mayhew M,
Mayhew M,
                                                                                                                                        Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
Al-Awgati Q,
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                                                                                                                                                                                                                                                   Disclosure; Page 33; 150pp; English
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an immune response,

WPI; 2001-663092/76

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                               cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response;
                                                                                                                                                                                                                                                                                                                     100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085;
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                                                                                                      The invention relates to a method of induction of
                                             protein and a melanoma antigen i.e. tyrosinase.
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                                                                        Disclosure, Page 23, 150pp; English
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LIVINGSTON P.
AL-AWQATI Q.
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Best Local Similarity
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising

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a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NVES01, MART antigens GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
HOE M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                               100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085;
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                                                   0; Mismatches
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72443 standard; peptide; 20 AA.
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase. Vyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980.AAU72481 represent melanoma antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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1 HWDFAWPW
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Homo sapiens.
Synthetic.
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Synthetic.
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(MAYH/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                      Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunothezapetic; hat shock protein; tyrosinase; BAGE; WTS801; GM2; tyrosinase related protein 1; tyrosinase related protein; yrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                            gp100-derived melanoma antigen, javelin peptide #13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein and a melanoma antigen i.e. tyrosinase
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26-FEB-2002 (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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RESULT 105

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen. Where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigens (AR, antigens, AR, antigens, AR, antigens, AR, antigens, AR, antigens, AR, antigens and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                   Anti cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; tive 0; Mismatches 0; Indels
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                                                                                              protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                         Disclosure; Page 26; 150pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU72294 standard; peptide; 20 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Houghton A, Livingston P,
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Best Local Similarity 100.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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                        WPI; 2001-663092/76
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                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock procein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTI980-AAUTI281 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; WYSB61; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                                                                                                                                                                                                    protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of
                                                                                                                                                                                     Livingston P, Al-Awgati Q,
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17-APR-2000; 2000US-0197462P
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                           HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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1 HWDFAWPW 8
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(MAYH/)
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             a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase protein 2, gp 100, MAGE antigens, BAGE antigens, NVES01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAV71980-AAV72481 represent melanoma antigen peptides of the invention
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comprising administration of an immunotherapeutic composition, comprising
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protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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(MAYH/)
(HOEM/)
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                                                           100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                  gp100-derived melanoma antigen, javelin peptide #42.
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Best Local Similarity 100.
Matches 8; Conservative
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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AAU72264;

RESULT 111

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAUT2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                                                                         Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunchkerapeutic; heat shock protein; tyrosinase; BAGE; VYSES01; GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                  gp100-derived melanoma antigen, javelin peptide #12
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANV1980-AAU72481 represent melanoma antigen peptides of the invention
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                17-APR-2000; 2000US-0197462P.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                      HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                            Anti cancer vaccine for the treatment of melanoma protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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(HOEM/) HOE M.
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Synthetic.
                    sapiens
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                                      Synthetic.
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(ALAW/)
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RESULT 114

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Sequence 20 AA;
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(ALAW/)
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                                                                                                        comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                           melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Melanoma antigen, MART-1, MAGE-1, gp100; cytostatic; immune response; immunotherapeutic; heat shock poteth; tyrosinase; BAGE; WYSB01, GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                 invention relates to a method of induction of an immune response,
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                                                                                                                                                                                                                                                      100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; ive 0; Mismatches 0; Indels
  Hoe
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  Mayhew M,
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                                           Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase.
  Al-Awgati Q,
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                                                                           Disclosure, Page 31; 150pp, English.
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                                                                                                                                                                                                                                                                                                                                                                             AAU72473 standard, peptide, 20 AA.
                                                                                                                                                                                                            antigen peptides of the invention
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  Livingston P,
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                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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(LIVI/) LIVINGSTON P.
(ALAW) AL-AWQATI Q.
(MAXYH/) MAYHEW M.
(HOEM/) HOE M.
                                                                                                                                                                                                                                                                                                   1 HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                  Sequence 20 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
Synthetic.
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  Houghton A,
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                         comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, Lyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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The invention relates to a method of induction of an immune response,
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72224 standard, peptide, 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprises a heat shock
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2;
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
davelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                    Gaps
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for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
antigen peptides of the invention
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                                                           4; Length 20;
                                                        100.0%; Score 64; DB 4; Length 20
100.0%; Pred. No. 0.0085;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                 Tyrosine-derived melanoma antigen, javelin peptide #28
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                                                                                                                                                                                         AAU72214 standard; peptide; 20 AA.
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                                                                                8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                         1 HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
                                                                    Similarity
                                                                                                                              1 HWDFAWPW
                                   Sequence 20 AA;
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(MAYH/)
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                                                                                                                                                                Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEB01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                             gp100-derived melanoma antigen, javelin peptide #3
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                                                     (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                     26-FEB-2002
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                                                                                                                                                                                                                                                                                                                                               Synthetic.
AAU72224;
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(LIVI/) 1
(ALAW/) 1
(MAYH/) 1
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Matches
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, related protein 1, tyrosinase, ratigence, BAGE antigens, NYESG1, MART antigens, GMZ, antigence portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                     Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYE801; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MART-1-derived melanoma antigen, javelin peptide #32.
                                                                                                                                                                                                                                                                                                                     Mayhew M,
                                                                                                                                                                                                                                                                                                                     Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 23; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU72393 standard; peptide; 20 AA.
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                Homo sapiens.
Synthetic.
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(HOEM/)
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                                                                                                 Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapettic; heat shock protein; tyrosinase; BAGE; NYBS01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                   MART-1-derived melanoma antigen, javelin peptide #3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein and a melanoma antigen i.e. tyrosinase
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                                  26-PEB-2002 (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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HWDFAWPW 20
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Best Local Similarity
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AMU71980-AMU72481 represent melanoma antigen peptides of the invention
                                                                                                    melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYBSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; favelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gp100-derived melanoma antigen, javelin peptide #83.
                      Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                       protein and a melanoma antigen i.e. tyrosinase
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                                                                                                  Anti cancer vaccine for the treatment of
                    Al-Awgati Q,
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                                                                                                                                                          Disclosure; Page 30; 150pp; English
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                    Houghton A, Livingston
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                          WPI; 2001-663092/76.
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                                                                                                                                                                                                                                                                                                                                                                                                  comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AMV11980-AAV72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                             Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                    Al-Awgati Q, Mayhew M, Hoe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WART-1-derived melanoma antigen, javelin peptide #33
                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 30; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAU72394 standard; peptide; 20 AA.
17-APR-2001; 2001WO-US012449
                                     17-APR-2000; 2000US-0197462P
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAXHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock procein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase ratigents, BAGE antigens, NYESO1, MART antigent of the fire portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTI281 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU72363 standard; peptide; 20 AA.
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                                                                                                                                                                                                                                                                                                           8, Conservative
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LIVINGSTON P.
AL-AWQATI Q.
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Matches 8; Conserv
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                                                                                                                                                                                                                                      Sequence 20 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-PEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-2001
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shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                        Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase abGE; WSES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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Best Local Similarity 100.0%; Pred. No. 0.0085;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                       100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.0085; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         gp100-derived melanoma antigen, javelin peptide #52.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 25; 150pp; English
                                                                                                                                                                                                                                                                                                     AAU72273 standard; peptide; 20 AA.
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                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2002 (first entry)
                                                                                                                           Best Local Similarity 100.
Matches 8; Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-663092/76
                                                                                                                                                                              1 HWDFAWPW 8
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                                                                        Sequence 20 AA;
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(HOEM/)
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RESULT 127

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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompactability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is human gpl00 MHC class I mutant peptide antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                            Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 100.0%; Score 64; DB 5; Length 20; Similarity 100.0%; Pred. No. 0.0085; 8; Conservative 0; Mismatches 0; Indels
                                                                                           1uman gp100 MHC class I mutant peptide antigen #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human gp100 MHC class I mutant peptide antigen #2.
                                                                                                                                                                                                                                                                                                                                                       /note= "MHC class I epitope"
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'note= "Javelin sequence"
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                                                                                                                                                                                                                                                                                                                  "Linker"
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                                                        (first entry)
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/note= "7
12.
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                                                                                                                                                                                                          domo sapiens
                                                    12-FEB-2002
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                   AAE13455;
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AAE13456
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AC AAE134
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DB Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of induction of an immune response, comprising administration of an immunotherspeutic composition. Comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigen, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AMJ71980-AAM72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                            Melanoma antigen, MART-1, MAGE-1, gp100; cytostatic, immune response; immunotherspeptic; heat shock protein; tyrosinase; BAGE; NYESO1, GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                       MAGE-3-derived melanoma antigen, javelin peptide #3.
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                                                                                                                                            AAU72474 standard; peptide; 20 AA
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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HWDFAWPW 20
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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RESULT 128 AAE13455 ID AAE134

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13-FEB-2003; 2003US-0447142P.
11-APR-2003; 2003US-0462469P.
                   WO2004071457-A2
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 AA;
                                                                                                                                                            Fletchner J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
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                                       26-AUG-2004
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  Synthetic
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                                                                                                                                                                      Barber B;
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Matches
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                                                                                                                                                                                                                                                                                                              A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; antimicrobial; cytostatic; vaccine; gene therapy; infectious disease; cancer.
         Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy; human.
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/note= "MHC class I epitope"
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                                                                                                                    13. .20
/note= "Javelin sequence"
                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                       Disclosure, Page 14; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADR69745 standard; peptide; 20 AA.
                                                                                                           "Linker"
                                                                                                                                                                                                                                                                        Hoe M;
                                                                                                                                                                                          17-APR-2001; 2001WO-US012567.
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                                                                                                                                                                                                                                                                        Mayhew M,
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                                                                                                                                                                                                                                  ROTHMAN J
                                                                                                                                                                                                                                          (MAYH/) MAYHEW M.
(HOEM/) HOE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 AA;
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                                                   Homo sapiens
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This invention relates to a novel hybrid antigen which comprises at least one antigenic domain of an infectious agent or tumour antigen and a binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                      Prince-Cohane K, Mehta S, Slusarewicz P, Andjelic S;
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                                                           13-FEB-2003; 2003US-0447142P.
11-APR-2003; 2003US-0462469P.
18-APR-2003; 2003US-0463746P.
16-SEP-2003; 2003US-0503417P.
13-FEB-2004; 2004WO-US004340.
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This invention relates to a novel hybrid antigen which comprises at least one antigenic domain of an infectious agent or tumour antigen and a binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
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shock protein-binding domains, useful for preventing or treating
infectious diseases or cancer.
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                                                                                                                                        Example 1, Page 38; 56pp, English
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11-APR-2003; 2003US-0462469P.
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16-SEP-2003; 2003US-0503417P.
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18-APR-2003; 2003US-0463746P.
16-SEP-2003; 2003US-0503417P.
12-FEB-2004; 2004US-00776521.
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18-APR-2003; 2003US-0463746P.
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gene therapy;
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invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
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100.0%; Score 64; DB 8; Length 20; 100.0%; Pred. No. 0.0085; rive 0; Mismatches 0; Indels
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ADU08486 standard; peptide; 20 AA.
      13-JAN-2005 (first entry)
         Hybrid antigen #1.
   ADU08486;
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Hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; hsp; immune response; infectious disease; cancer; cytostatic; antimicrobial; immunostimulant.

Synthetic. Unidentified

WO2004091493-A2.

28-OCT-2004

09-APR-2004; 2004WO-US010983.

2003US-0463746P. 2003US-0503417P. 2004US-00776521. 11-APR-2003; 8-APR-2003; 16-SEP-2003;

(MOJA-) MOJAVE THERAPEUTICS INC 2004US-00820067. 13-FEB-2004;

2004WO-US004340

Andjelic S; Slusaregicz P, Mehta S, Prince-Cohane K, Flechtner JB, Barber BH;

WPI; 2004-775516/76.

Hybrid antigen useful for treating an infectious disease or cancer, comprises an antigenic domain from the infectious agent or cancer joined to a heat shock protein binding domain through an improved linker peptide

Example 2; Page 59; 99pp; English

The invention relates to hybrid antigens comprising at least one antigenic domain of an infectious agent or tumour antigen, at least one binding domain that non-covalently binds to a heat shock protein (hsp), and at least one peptide linker between them. Also disclosed are: (a) a composition for inducing an immune response to an infectious agent or tumour antigen comprising at least one of the hybrid antigens or a complex of at least one heat shock protein and at least one of the hybrid antigens, (b) a method for inducing an immune response to an infectious agent or tumour antigen by administering a hybrid antigen and a heat shock protein, where the hybrid antigens and the heat shock protein are non-covalently bound, and (c) treating an infectious disease or cancer by

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administering a hybrid antigen and a heat shock protein. The heat shock protein is preferably hep70. The composition is administered via oral or parenteral route. The hybrid antigen is useful in preparing a composition for treating or preventing cancer or infectious disease. The new peptide linkers give the antigens improved activity. Note: Many of the SEQ ID Nos are replicated more than once in the specification but the sequences of these replicated SEQ ID Nos are not the same. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    melanoma comprises a heat shock
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100.0%; Pred. No. 0.0085;
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                                                                                                                                                                                                                                   Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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                                                                                                                                                                                                             Local Similarity
                                                                                                                                                               Sequence 20 AA;
                                                                                                                              hybrid antigen
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, ratigens of the selected protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                   Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                  gp100-derived melanoma antigen, javelin peptide #102.
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                                                                              AAU72323 standard; peptide; 21 AA.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72323
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                                     100.0%; Score 64; DB 4; Length 21; 100.0%; Pred. No. 0.0089;
                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                             gp100-derived melanoma antigen, javelin peptide #133.
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                                                                            0; Mismatches
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nes 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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comprising administration of an immunotherapeutic composition, Comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEG1, MART antigens, GM2, antigent cortions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                          melanoma comprises a heat shock
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                       Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunctherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gp100-derived melanoma antigen, javelin peptide #112.
                 gp100-derived melanoma antigen, javelin peptide #103
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protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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RESULT 139

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                              treatment of melanoma comprises a heat shock
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                                                                                Disclosure; Page 31; 150pp; English
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(HOEM/)
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antigen peptides of the invention

RESULT 145

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase entelated protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                                             Melanoma antigen, MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; NYESE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                        MART-1-derived melanoma antigen, javelin peptide #22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Al-Awgati Q,
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26-FEB-2002 (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEG1, MART antigens, GN2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                    Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response;
immunotherapeutic; heat shock protein; tyrosinase; BAGE; WYSBG11, GM2;
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                               gp100-derived melanoma antigen, javelin peptide #132.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a melanoma antigen i.e. tyrosinase
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                                                                                                       AAU72353 standard; peptide; 21 AA.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
HOE M.
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8; Conserva
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HWDFAWPW
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(MAYH/)
(HOEM/)
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RESULT 146

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AAU72383 ID AAU XX AC AAU XX

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                   Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYE801; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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17-APR-2000; 2000US-0197462P.
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nes 8; Conservative
                                        (HOUG/) HOUGHTON A.
(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(ALAW/) AL-AWQ
(MAYH/) MAYHEW
(HOEM/) HOE M.
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(MAYH/)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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      Homo sapiens.
Synthetic.
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RESULT 148

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, WART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the had shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS81 represent melanoma antigen peptides of the invention

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Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                    vaccine for the treatment of melanoma comprises a heat shock
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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protein and a melanoma antigen i.e. tyrosinase
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Length 21; 0; Indels

Query Match
100.0%; Score 64; DB 4;
Best Local Similarity 100.0%; Pred. No. 0.0089;
Matches 8; Conservative 0; Mismatches 0;

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Melanoma antigen, MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human. MAGE-1-derived melanoma antigen, javelin peptide #32. AAU72433 standard; peptide; 21 26-FEB-2002 (first entry) AAU72433; (/SDOH) HOEM/) (LIVI/) (ALAW/) MAYH/ RESULT 151 AAU72433 Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSB61; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human. MAGE-1-derived melanoma antigen, javelin peptide #23.

The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma Anti cancer vaccine for the treatment of melanoma comprises a heat shock Hoe M; Mayhew M, protein and a melanoma antigen i.e. tyrosinase Al-Awgati Q, Disclosure; Page 32; 150pp; English 17-APR-2001; 2001WO-US012449. 17-APR-2000; 2000US-0197462P Houghton A, Livingston P, HOUGHTON A. LIVINGSTON P. AL-AWQATI Q. MAYHEW M. WPI; 2001-663092/76. WO200178655-A2. Homo sapiens. 25-OCT-2001 Synthetic.

Hoe M;

Mayhew M,

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antigen peptides of the invention

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AAU72373 standard; peptide; 21 AA.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, MAGE antigens, MA, antigens, MA, antigens, MA, antigens, MA, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                 MART-1-derived melanoma antigen, javelin peptide #12.
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protein and a melanoma antigen i.e. tyrosinase
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAXH/) MAXHEW M.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma shigen is covalently bound to a javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUT1980-AAUT2481 represent melanoma antigen peptides of the invention
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                                                                                    Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; hat shock protein; tyrosinase; BAGE; NYESO1; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                   MART-1-derived melanoma antigen, javelin peptide #13.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen; where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma.

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protein and a melanoma antigen i.e. tyrosinase.
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17-APR-2001; 2001WO-US012449

(LIVI/) (ALAW/) (MAYH/)

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, pacted from tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                             melanoma comprises a heat shock
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                                                                                                                                                                                 The invention relates to a method of induction of an immune response,
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tive 0; Mismatches 0; Indels
Hoe
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Mayhew M,
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                                                                             Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase
Al-Awgati Q,
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                                                                                                                                           Disclosure; Page 33; 150pp; English
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Houghton A, Livingston P,
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 HWDFAWPW 21
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                   Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWOATI Q.
MAYHEW M.
HOE M.
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            comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, sp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule; where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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The invention relates to a method of induction of an immune response
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
HOE M.
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for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                  gp100-derived melanoma antigen, javelin peptide #113.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, MAGE antigens, MA, antigens, MA, antigens, MA, antigens of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                      Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; WSES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hoe
                                                                gp100-derived melanoma antigen, javelin peptide #41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Houghton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 25; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU72258 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                     17-APR-2001; 2001WO-US012449.
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                    26-FEB-2002 (first entry)
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 AA;
                                                                                                                                                                                                                                                                                  WO200178655-A2
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                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel hybrid antigen which comprises at least one antigenic domain of an infectious agent or tumour antigen and a binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a peptide which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                        hybrid antigen; antigenic domain; infectious agent; tumour antigen; binding domain; heat shock protein; antimicrobial; cytostatic; vaccine; gene therapy; infectious disease; cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New hybrid antigens comprising an antigenic domain and improved heat shock protein-binding domains, useful for preventing or treating infectious diseases or cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fletchner J, Prince-Cohane K, Mehta S, Slusarewicz P,
                                                                                                                                                                                                                                                                                Novel hybrid antigen-related peptide #1333.
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                                                                                                                                                 ADR69753 standard; peptide; 21 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 38; 56pp; English
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18-APR-2003; 2003US-0463746P.
16-ESB-2003; 2003US-053417P.
12-FSB-2004; 2004US-00776521.
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HWDFAWPW 21
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Matches 8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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Barber B;

AAU72262;

SXS

RESULT 162

Query Match

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Synthetic.

ADR69753;

RESULT 161

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ADR69753

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Gaps

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17-APR-2000; 2000US-0197462P
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                             HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                            WPI; 2001-663092/76
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                                                                                                                                                                                                                                                                                                            Sequence 25 AA;
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Synthetic.
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                            (HOUG/)
(LIVI/)
(ALAW/)
(MAYH/)
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(HOEM/)
                                                                     HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANU71980-AAU712481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                          Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                   Hoe M;
                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 4; Length 24; 100.0%; Pred. No. 0.01;
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                                                                                                                                                                                    Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                   Houghton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                          Disclosure; Page 25; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU72402 standard; peptide; 25 AA
                                                                               17-APR-2001; 2001WO-US012449.
                                                                                                  17-APR-2000; 2000US-0197462P.
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                                                                                                                      HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
                                                                                                                                                                                                     WPI; 2001-663092/76.
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Best Local Similarity
                                                                                                                                                       MAYHEW M.
                                       WO200178655-A2.
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24 AA;
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                                                                                                                                                               (HOEM/) HOE M
           sapiens.
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Synthetic.
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                                                           25-OCT-2001
                   Synthetic.
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           Homo
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javalin molecule, where the melanoma antigen bound to the javalin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                         Hoe M;
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                                                                                                                                                                                                                                                                                                                                                                                         Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                         Houghton A, Livingston P, Al-Awgati Q,
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17-APR-2000; 2000US-0197462P.
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                                                                                                                                                    Best Local Similarity
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                                                                                                                       Sequence 25 AA;
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                                                                                                                                           Query Match
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(LIVI/)
(ALAW/)
(MAYH/)
(HOEM/)
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                                                                                     The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, more the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                      Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                               Melanoma antigen, MART-1, MAGE-1, gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGB; NYSB01, GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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Hoe
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Mayhew M,
                                                 a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer vaccine for the treatment of
Al-Awgati O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Al-Awgati Q,
                                                                     Disclosure; Page 25; 150pp; English.
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                                                                                                                                                                                           antigen peptides of the invention
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Livingston P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Houghton A, Livingston P,
                                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                                                                                                                                                                                                                        8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                             1 HWDFAWPW
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                                                                                                                                                                                                                Sequence 25 AA;
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                                                protein and
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Houghton A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                          comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, tyrosinase protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEG1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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The invention relates to a method of induction of an immune response,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72189 standard; peptide; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                      immunotherapeutic, heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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100.0%; Pred. No. 0.011;
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                                                                                                                                                              AAU72452 standard; peptide; 25 AA.
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                                                                       8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                 Query Match
Best Local Similarity
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                               Sequence 25 AA;
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigens antigens, GMZ, antigens, GMZ, antigens, Dortions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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                                                                                                                          Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSB01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                   Tyrosine-derived melanoma antigen, javelin peptide #3.
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LIVINGSTON P.
AL-AWQATI Q.
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Best Local Similarity
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100.0%; Score 64; DB 4; Length 25; larity 100.0%; Pred. No. 0.011; Conservative 0; Mismatches 0; Indels

Best Local Similarity Matches 8; Conserv

Query Match

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigens, DAGE antigens, BAGE antigens of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                            cancer vaccine for the treatment of melanoma comprises a heat shock
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javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                              protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                             (HOUG/) HOUGHTON A.
(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(HOEM/) HOE M.
                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapettic; heat shock protein; tyrosinase; BAGE; WYSB61; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; favelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                MART-1-derived melanoma antigen, javelin peptide #7.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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                                26-FEB-2002
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   AAU72368;
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(HOEM/)
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, BAGE antigens, NYBs01, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-ANUTIS481 represent melanoma antigen peptides of the invention
                                                                                         melanoma comprises a heat shock
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                   Hoe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gp100-derived melanoma antigen, javelin peptide #91.
                 Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein and a melanoma antigen i.e. tyrosinase.
                                                                                       Anti cancer vaccine for the treatment of melanc
protein and a melanoma antigen i.e. tyrosinase
                 Al-Awgati Q,
                                                                                                                                                  Disclosure; Page 26; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU72312 standard; peptide; 25 AA.
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                 Livingston P,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                    WPI; 2001-663092/76.
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1 HWDFAWPW
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                   Houghton A,
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(LIVI/)
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                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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17-APR-2001; 2001WO-US012449
                                     17-APR-2000; 2000US-0197462P
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                                                                        HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention

Sequence 25 AA;

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             The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GM2, antigence portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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Matches 8, Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
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related protein 2, gp 100, MAGE antigens, BAGE antigens, NYBS01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                Melanoma antigen, MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; WYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2;
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                                                                                             gp100-derived melanoma antigen, javelin peptide #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 24; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                           17-APR-2001; 2001WO-US012449
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                                                       (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                                                                                                      Homo sapiens
                                                       26-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock proctein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase ralated protein 2, gp 100, MAGE antigens, BAGE antigens, NYE801, MART antigen, GMZ, antigent cortions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat for the treatment of melanoma to the treatment of melanoma ANT/1980-AAUT/2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tvrosinase.
                                                                                                                                                                                                                                                                                                          Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                       Jp100-derived melanoma antigen, javelin peptide #81
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                                                                                                                                                 AAU72302 standard; peptide; 25 AA
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72238 ID AAU7

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigens, GMZ, antigens, GMZ antigens, BAGE antigens of these: The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                   melanoma comprises a heat shock
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 4; Length 25; 100.0%; Pred. No. 0.011; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                Houghton A, Livingston P, Al-Awgati Q, Mayhew M,
                                                                                                                                                                                                                                                                                                                 Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase.
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                 17-APR-2001; 2001WO-US012449.
                                                             17-APR-2000; 2000US-0197462P.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                     HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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(LIVI/) LIVINGS
(ALAW/) AL-AWQA
(MAYH/) MAYHEW
(HOEM/) HOE M.
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Synthetic.
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  tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                             Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                           Livingston P, Al-Awgati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Disclosure; Page 31; 150pp; English.
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                                                                                                                       comparising administration of an immunotherapeutic composition, comparising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from trosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESU, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANV1980-AAV72481 represent melanoma antigen peptides of the invention
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                                                            comprises a heat shock
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                                                                                                              The invention relates to a method of induction of an immune response
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                    Hoe M;
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                    Mayhew M,
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                                                             melanoma
                                                           Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase
                   Livingston P, Al-Awgati Q,
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                                                                                            Disclosure; Page 30; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                AAU72422 standard; peptide; 25 AA.
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Best Local Similarity 100.vv
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                        WPI; 2001-663092/76.
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Synthetic.
(HOEM/) HOE
                    Houghton A,
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase artigens, MYESO1, MART antigens GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS481 represent melanoma antigen peptides of the invention
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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antigen bound to the javelin molecule is non-covalently bound to the heat shock protestn. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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MAYHEW M.
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RESULT 186

AAU72190

Score 64; DB 4; Length 25; Pred. No. 0.011;

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Query Match Best Local Similarity

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEGO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEsOl; GM2;
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
javelin molecule; melanoma antigen recognised by T cells-1; human.
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                        Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; VYSESU1; GN2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                    Pyrosine-derived melanoma antigen, javelin peptide #4
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      AAU72190 standard; peptide; 25
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, MNEs01, MART antigens, GMY, antigens, GMY, antigens, Oportions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                        Anti cancer vaccine for the treatment of melanoma comprises a heat shock
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                                                             Al-Awgati Q,
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                                                                                                                                                                                                      Disclosure; Page 24; 150pp; English
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(MAYH/) MAYHEW M.
(HOEM/) HOE M.
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                                                                                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                      17-APR-2001; 2001WO-US012449
                                                                              17-APR-2000; 2000US-0197462P.
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LIVINGSTON P.
AL-AWQATI Q.
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LIVINGSTON P.
AL-AWQATI Q.
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RESULT 189

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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESU, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention relates to a method of induction of an immune Disclosure; Page 29; 150pp; English

Sequence 25 AA;

Gaps ö 0; Indels 100.0%; Score 64; DB 4; Length 100.0%; Pred. No. 0.011; 0; Mismatches 8; Conservative Best Local Similarity Query Match Matches

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1 HWDFAWPW

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AAU72418 standard; peptide; 25 AA.

(first entry) 26-FEB-2002 AAU72418;

MAGE-1-derived melanoma antigen, javelin peptide #17.

Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immuncherapeutic; haat shock protein; tyrosinase; BAGE; NYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T celle-1; human.

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17-APR-2001; 2001WO-US012449

17-APR-2000; 2000US-0197462P

HOUGHTON A. LIVINGSTON P. AL-AWQATI Q. (HOUG/) (LIVI/) (ALAW/)

MAYHEW M. HOE M MAYH/) (HOEM/)

Ξ Mayhew M, Al-Awgati Q, Houghton A, Livingston P,

WPI; 2001-663092/76.

comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen as elected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, 9p 100, MAGE antigens, MAGE antigens, MAGE antigens, MAGE antigens, CM2, antignic portions and combinations of these. The melanoma Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase. The invention relates to a method of induction of an immune response Disclosure; Page 31; 150pp; English AAU72418

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ö antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention where the melanoma Gaps ö y maccn 100.0%; Score 64; DB 4; Length 25; Local Similarity 100.0%; Pred. No. 0.011; hes 8; Conservative 0; Mismarches antigen is covalently bound to a javelin molecule, 1 HWDFAWPW 8 HWDFAWPW Ż Sequence 25 Query Match Best Loca Matches 88888888 셤 à

gp100-derived melanoma antigen, javelin peptide #57. (first entry) 26-FEB-2002

AAU72278 standard; peptide; 25 AA.

RESULT 192

AAU72278

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AAU72278

Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.

sapiens Synthetic.

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25-OCT-2001

17-APR-2001; 2001WO-US012449.

17-APR-2000; 2000US-0197462P. HOUGHTON A. LIVINGSTON P. AL-AWQATI Q. MAYHEW M. HOUG/) (LIVI/) (ALAW/) (MAYH/)

HOEM/)

Mayhew M, P, Al-Awgati Q, Houghton A, Livingston

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WPI; 2001-663092/76

Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.

Disclosure; Page 25; 150pp; English

The invention relates to a method of induction of an immune response, comprising administration of an immunocherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase arelated protein 1, tyrosinase antigens for the portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention

Sequence 25 AA;

Query Match

DB 4; 100.0%; Score 64;

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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; NYBS01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                     gp100-derived melanoma antigen, javelin peptide #1.
            AAU72222 standard; peptide; 25
                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens BAGE antigens, NYESO1, MART antigens, GM2, antigenfc portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
             Gaps
                                                                                                                                                                                                                               Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immuncherapeutic; heat shock protein; tyrosinase; BAGE; NYSEG1; GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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Pred. No. 0.011; ; Mismatches 0; Indels
                                                                                                                                                                                                         MAGE-1-derived melanoma antigen, javelin peptide #11.
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                                                                                                                              AAU72412 standard, peptide; 25 AA.
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100.0%;
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                                                                                                                                                                                 (first entry)
              Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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                                                                                                    RESULT 193
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, in the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU11980-AAU72481 represent melanoma antigen peptides of the invention
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17-APR-2001; 2001WO-US012449.
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                                                                                                          (HOUG/) HOUGHTON A.
(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
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RESULT 194

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                    Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; thmunocherapeutic; heat shock protein; tyrosinase; BAGE; WYSE01; GMZ; YYSOsinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein and a melanoma antigen i.e. tyrosinase
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                     melanoma comprises a heat shock
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protein and a melanoma antigen i.e. tyrosinase
                                                          P, Al-Awgati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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MAYHEW M.
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(ALAW/) (MAYH/) (HOEM/) )
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                                                                            comprising administration of an immunotherapeutic composition, Comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, NART antigens GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                The invention relates to a method of induction of an immune response
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                                                                                                                                                                                                                                                                                                        100.0%; Score 64; DB 4; Length 26; 100.0%; Pred. No. 0.011; ative 0; Mismatches 0; Indels
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protein and a melanoma antigen i.e. tyrosinase
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                                Disclosure; Page 27; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU72348 standard, peptide; 26
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
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        antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protedin. The composition is useful for inducing an immune response for the treatment of melanoma. AMU71980-AMU72481 represent melanoma antigen peptides of the invention
GM2, antigenic portions and combinations of these. The melanoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 27; 150pp; English
                                                                                                                                                                                                                                                                                                                        AAU72322 standard; peptide; 26 AA
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                                                                                                                                                                          Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                           Sequence 26 AA;
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 antigens,
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Best Local
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Sequence 26 AA,

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                             MART-1-derived melanoma antigen, javelin peptide #17.
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                                                                                      AAU72378 standard; peptide; 26 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer vaccine for the
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                                                                                                                                               (first entry)
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) AL-AWQATI Q.
MAYHEW M.
HOE M
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                              Homo sapiens.
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                                                                                                     Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                  MART-1-derived melanoma antigen, javelin peptide #21.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 29; 150pp; English
                    AAU72382 standard; peptide; 26 AA
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                                                           WO200178655-A2
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                                                                                                                                                            Homo sapiens.
Synthetic.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens BAGE antigens, NYEsO1, MART antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTISBO-ANUT2481 represent melanoma antigen peptides of the invention
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; NYBE01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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(HOUG/) (LIVI/) (ALAW/) (MAYH/) (HOEM/)

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigens, GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                            melanoma comprises a heat shock
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                                                                                                Mayhew M,
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                                                                                                Al-Awgati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
LIVINGSTON P.
                  AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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WO200178655-A2
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cancer vaccine for the treatment of melanoma comprises a heat shock
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protein and a melanoma antigen i.e. tyrosinase.
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                                                Disclosure; Page 34; 150pp; English
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LIVINGSTON P.
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MAYHEW M.
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related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the hack shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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(LIVI/) LIVINGSTON P.
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(MAYH/) MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU72248 standard; peptide; 26 AA.
                                                          AAU72338 standard; peptide; 26 AA.
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                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-663092/76
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                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                        RESULT 210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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        100.0%; Score 64; DB 4; Length 26; larity 100.0%; Pred. No. 0.011; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                              gp100-derived melanoma antigen, javelin peptide #31
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                                                                                                                                                                                                                                        AAU72252 standard; peptide; 26 AA.
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AL-AWQATI Q.
MAYHEW M.
Query Match
Best Local Similarity
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17-APR-2000; 2000US-0197462P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunothezapettic; heat shock protein; tyrosinase; BAGE; NYRSB01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapetic; heat shock protein; tyrosinase; BAGE; NYESO1; GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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gp100-derived melanoma antigen, javelin peptide #27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein and a melanoma antigen i.e. tyrosinase.
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les 8; Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWOATI Q.
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Matches
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, related protein 1, tyrosinase antigens, BAGE antigens, NYESG1, MART antigens, GM2, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTIS481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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17-APR-2001; 2001WO-US012449.
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                                                                                                                                             (HOUG/) HOUGHTON A. (LIVI/) LIVINGSTON P. (ALAW) AL-AWQATI Q. (MAYH/) MAYHEW M.
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HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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                                                                                                                                                        comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                 Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                  The invention relates to a method of induction of an immune response
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                                                           Mayhew M,
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                                                           Al-Awgati Q,
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                                                                                                                              Disclosure; Page 27; 150pp; English
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                                                           Houghton A, Livingston P,
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 HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
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melanoma comprises a heat shock
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    Anti cancer vaccine for the treatment of melanc
protein and a melanoma antigen i.e. tyrosinase.
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is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, some the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GMZ, antigens, Gportions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                AAU72318 standard; peptide; 26 AA.
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase abdGE; NYBS01; GM2; Yyxosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
             WAGE-1/3-derived melanoma antigen, javelin peptide #21
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protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                               Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunchezpetic; hat shock protein; tyrosinase; BAGE; NYSS61; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of induction of an immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 28; 150pp; English
                                                       AAU72352 standard; peptide; 26 AA
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                                                                                                                                                                                                                                                                                                                                                                       17-APR-2001; 2001WO-US012449
                                                                                                                                                                                                                                                                                                                                                                                                      17-APR-2000; 2000US-0197462P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Houghton A, Livingston P,
                                                                                                                   (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                                                                                                                                                                                                                                                         WO200178655-A2
                                                                                                                                                                                                                                                               sapiens
                                                                                                                   26-FEB-2002
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                                                                                                                                                                                                                                                                              Synthetic
                                                                                      AAU72352
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(ALAW/)
(MAYH/)
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                        RESULT 218
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ID AAU7
XX
AC AAU7
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DT 26-F
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melanoma comprises a heat shock

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Mayhew M,

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                                                                                                                                      comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Melanoma antigen; MART-1; MAGE-1; gpl00; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                            The invention relates to a method of induction of an immune response,
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Disclosure, Page 33; 150pp; English.
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(HOUG/) (ALAW/) (HOEM/)

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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by at etchering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken MHC class II peptide antigen
                                                                                                                                                                                                                                                                                   A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
100.0%; Score 64; DB 5; Length 27;
Best Local Similarity 100.0%; Pred. No. 0.012;
Matches 8; Conservative 0; Mismatches 0; Indels
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/note= "Javelin sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chicken MHC class II peptide antigen #2
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                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 13; 47pp; English.
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                                                                                                                                                                                                             Hoe M;
                                                      17-APR-2001; 2001WO-US012567.
                                                                                          17-APR-2000; 2000US-0197462P.
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/note= "'
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/note=
                                                                                                                                                                                                             Rothman JE, Mayhew M,
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                                                                                                                            (ROTH/) ROTHMAN J E. (MAYH/) MAYHEW M. (HOEM/) HOE M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 27 AA;
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                25-OCT-2001.
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Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response;
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                                                                                                                                                                                                                                                                                                           Hoe M;
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                                                                                                                                                                                                                                                                                                           loughton A, Livingston P, Al-Awgati Q, Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                        protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "MHC class II epitope"
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/note= "Javelin sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chicken MHC class II peptide antigen #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 33; 150pp; English
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                                                                                                              17-APR-2001; 2001WO-US012449.
                                                                                                                                                  17-APR-2000; 2000US-0197462P
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                                                                                                                                                                                       HOUGHTON A.
LIVINGSTON P.
AL-AWQAII Q.
MAYHEW M.
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malignancy; chicken.
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tes 8; Conserv
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Synthetic.
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Query Match

Best Loc Matches

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RESULT 221 AAE13449 Key Region Region Region

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Gaps

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protein and a melanoma antigen i.e. tyrosinase
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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                                                                                                                                                                                     Sequence 30 AA;
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                                                                                                                                                                     The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major is stocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken MHC class II peptide antigen
                                                                                                   A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYSS01; GMZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                                                Disclosure, Page 13; 47pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU72255 standard; peptide; 30 AA.
                                                       Hoe M;
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Matches 8; Conservative
                                                      Rothman JE, Mayhew M,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                             NPI; 2002-017594/02
          ROTHIMAIN J
                     МАҮНЕМ М.
НОЕ М.
                                                                                                                                                                                                                                                                                         Sequence 27 AA;
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(MAYH/)
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GMZ, antigen for portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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protein and a melanoma antigen i.e. tyrosinase.
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Disclosure; Page 25; 150pp; English
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histocompatability complex class I (MHC) and the other MHC class II. The santigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken MHC class I peptide antigen
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 antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                                                                  Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy; chicken.
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/note= "Javelin sequence"
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                                                                                                                                                                                                                                                                                 Chicken MHC class I peptide antigen #3
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                                                                                                                                                                                                                  AAE13448 standard; peptide; 30 AA.
                                                     antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             new antigenic complex comprising epitopes non-covalently joined to a sat shock protein by a molecular tether designated a javelin are useful b treat or prevent infectious disease or malignancy.
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                                                                                                                                                                                                                                                                                                                             AAE13454 standard; peptide; 30 AA.
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'note= "Linker"
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treatment or prevention of infectious diseases and malignancies. The present sequence is Herpes simplex virus MHC class I peptide antigen
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                                                                              100.0%; Score 64; DB 5; Length 30; 100.0%; Pred. No. 0.013; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                              AAU72295 standard; peptide; 31 AA
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LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                  Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; haat shock protein; tyrosinase; BAGE; WYSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                     Tyrosine-derived melanoma antigen, javelin peptide #6.
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                            AAU72192 standard; peptide; 31 AA.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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RESULT 228
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigens, DW4G antigens, BAGE antigens, NYE801, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                          Al-Awgati Q,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                        Homo sapiens.
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                                                                                                                       Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javalin molecule; melanoma antigen recognised by T celle-1; human.
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tive 0; Mismatches 0; Indels
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                                                                                   gp100-derived melanoma antigen, javelin peptide #9.
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LIVINGSTON P.
AL-AWQATI Q.
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Best Local Similarity
Matches 8; Conserv
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Disclosure; Page 24; 150pp; English
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            Houghton A, Livingston P,
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(LIVI/) LIVINGSTON P
(ALAW/) AL-AWQATI Q.
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                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                       Mayhew M,
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                                                                                                     Livingston P, Al-Awgati Q,
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                    17-APR-2000; 2000US-0197462P
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Matches 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                                      MAYHEW M.
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                                                                                                                                                                            melanoma comprises a heat shock
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    Mayhew M,
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    Al-Awgati Q,
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigen 5 MAGE antigens, MART antigen, GMY, antigenfc portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUT1980-ANUT2481 represent melanoma antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                  AAU72400 standard; peptide; 31 AA.
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Best Local Similarity 100.
Matches 8; Conservative
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(MAYH/) MAYHEW
(HOEM/) HOE M.
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MAGE-1/3-derived melanoma antigen, javelin peptide #4.
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                                                                                                                                          WAGE-1-derived melanoma antigen, javelin peptide #4.
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                                                                              AAU72405 standard; peptide; 31 AA
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase, tyrosinase related protein 1, tyrosinase antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma antigen peptides of the invention
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase sABGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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AAU72445 standard; peptide; 31 AA.

RESULT 237

AAU72445 ID AAU7

melanoma comprises a heat shock

Mayhew M,

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigens portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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           17-APR-2001; 2001WO-US012449.
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Matches 8; Conservative
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                                                                                      HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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LIVINGSTON P.
AL-AWQATI Q.
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(MAYH/) MAYHEW
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                                                       Homo sapiens
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comportaing administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from trosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYE801, MART antigens GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                           melanoma comprises a heat shock
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                                                                                                                                                                         The invention relates to a method of induction of an immune response.
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                                Mayhew M,
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                                                                                          Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase
                                Al-Awgati Q,
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                                                                                                                                        Disclosure; Page 23; 150pp; English.
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                                Livingston P,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                              The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, DAGE antigens, BAGE antigens, NRSEOL, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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Disclosure; Page 25; 150pp; English
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYE901, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule, ann-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANU1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                 MAGE-1-derived melanoma antigen, javelin peptide #9.
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     Mismatches
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     Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Synthetic.
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AAU72305
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antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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100.0%; Pred. No. 0.013;
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LIVINGSTON P.
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEsOl; GM2;
tyrosinase related protein 1; tyrosinase related protein 2; vaccine;
javelin molecule; melanoma antigen recognised by T cells-1; human.
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AL-AWQATI Q.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 31 AA;
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                                                                                                             WO200178655-A2
                                                                                                                                                                                                                                                                                                        HOE M.
                                                                 Homo sapiens.
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                                                                                Synthetic.
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(ALAW/)
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, pp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                            Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinashase; BAGE; NYBS01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method of induction of an immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hoe M;
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                                                                                               FP100-derived melanoma antigen, javelin peptide #84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gp100-derived melanoma antigen, javelin peptide #79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Houghton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 26; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU72300 standard; peptide; 31 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antigen peptides of the invention
    AAU72305 standard, peptide, 31
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                                                                 (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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Synthetic.
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                                  AAU72305
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(LIVI/)
(ALAW/)
(MAYH/)
(HOEM/)
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comprises a heat shock

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Mayhew M,

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Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                                   Houghton A, Livingston P,
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Best Local Similarity 100...
... 8; Conservative
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(MAYH/) MAYHEW M.
(HOEM/) HOE M.
                                                                                                                                                                                                                                                                                 Sequence 31 AA;
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Synthetic.
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(MAYH/)
(HOEM/)
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                                                                                                                                                                                                                                                        comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens protein 2, gp 100, MAGE antigens, BAGE antigens, NYEs01, MAKT antigens, GMZ, antigent portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma sintigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANV11980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                      melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; WSES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAGE-1/3-derived melanoma antigen, javelin peptide #9
                                                                                                                                          Houghton A, Livingston P, Al-Awgati Q, Mayhew M,
                                                                                                                                                                                       Anti cancer vaccine for the treatment of meland protein and a melanoma antigen i.e. tyrosinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU72450 standard; peptide; 31 AA.
                       17-APR-2001; 2001WO-US012449
                                              17-APR-2000; 2000US-0197462P
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                                                                     HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
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                                                                                                                                                               WPI; 2001-663092/76
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                                                                                                       MAYHEW M.
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Synthetic.
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  25-OCT-2001
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(HOEM/)
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(ALAW/)
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AAU72450
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                                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase BAGE; WSEB01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vacCine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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    Hoe
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Mayhew M,
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                                                                                                                                                                                                                     protein and a melanoma antigen i.e. tyrosinase
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    Al-Awgati Q,
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Synthetic.
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Query Match
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antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUTI980-AAUT2481 represent melanoma antigen peptides of the invention
                                                                                   Sequence 31 AA;
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                                                                                                            Query Match
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                                                      comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AU71980-AAU72481 represent melanoma antigen peptides of the invention
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                                           to a method of induction of an immune
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                                                                                                                                                                                                                            100.0%; Score 64; DB 4;
100.0%; Pred. No. 0.013;
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                  Disclosure, Page 23; 150pp, English.
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                                                                                                                                                                                                                                                                                                                                                                       AAU72365 standard; peptide; 31 AA
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                                                                                                                                                                                                                                                       8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                           The invention relates
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                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                               1 HWDFAWPW
                                                                                                                                                                                                    Sequence 31 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase ARGE; NYES01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gpl00-derived melanoma antigen, javelin peptide
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                                                               0; Mismatches
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                                                               8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                              Local Similarity
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100.0%; Score 64; DB 4; Length 31;

(first entry)

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AAU72280 standard; peptide; 31 AA.
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                                           AAU72280;
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(HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                      Gaps
                                                                                                                                                                                                              Melanoma antigen, MART-1; MAGE-1; gp100; cytostatic; immune response; immunothergapettic; heat shock protein; tyrosinase; BAGE; WYSE01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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100.0%; Pred. No. 0.013;
ive 0; Mismatches 0; Indels
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                      Indels
                                                                                                                                                                                          gp100-derived melanoma antigen, javelin peptide #54.
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Pred. No. 0.013;
                     0; Mismatches
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                                                                                                                       AAU72275 standard; peptide; 31 AA.
         100.04;
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                                                                                                                                                                   (first entry)
                      Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-663092/76
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Best Local Similarity
Matches 8; Conserv
           Best Local Similarity
                                          1 HWDFAWPW
                                                                HWDFAWPW
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                                                                                                                                                                                                                                                                      Homo sapiens.
Synthetic.
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(HOEM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
                                                 Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; WRSB01; GM2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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gp100-derived melanoma antigen, javelin peptide #59.
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Nes 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HOUG/) HOUGHTON A. (LIVI/) LIVINGSTON P. (ALAW/) AL-AWQATI Q.
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1 HWDFAWPW HWDFAWPW

RESULT 253

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Synthetic
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                                                                         (HOUG/)
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                                                                                                                                                                                                                                                                               Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYBS01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                             17-APR-2001; 2001WO-US012449
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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                                                     sapiens.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GM2, antigen fortions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAUT2481 represent melanoma antigen peptides of the invention
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                                                                17-APR-2001; 2001WO-US012449.
                                                                                                                                 17-APR-2000; 2000US-0197462P.
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nes 8; Conservative
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) AL-AWQATI Q.
MAYHEW M.
HOE M
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LIVINGSTON P.
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protein and a melanoma antigen i.e. tyrosinase
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                                                                                                                            The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigen for the antigen and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANUTIS80-AAUTI281 represent melanoma antigen peptides of the invention
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                                                                               comprises a heat shock
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                                         Hoe M;
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                                         Mayhew M,
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                                                                              Anti cancer vaccine for the treatment of melanoma protein and a melanoma antigen i.e. tyrosinase.
                                         Al-Awgati Q,
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                                                                                                             Disclosure, Page 24; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     AAU72270 standard; peptide; 31 AA.
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                                         Livingston P,
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
 AL-AWQATI Q.
MAYHEW M.
                                                            WPI; 2001-663092/76
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Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW
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                                                                                                                                                                                                                                                       Sequence 31 AA;
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                   HOE M
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                                         Houghton A,
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(ALAW/)
(MAYH/)
(HOEM/)
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(MAYH/)
(HOEM/)
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                                          The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, tyrosinase related protein 1, tyrosinase antigens, 6MZ, antigens, 6MZ, antigens antigens of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AdVI1980-AAVI2481 represent melanoma antigen peptides of the invention
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heat shock protein by a molecular tether designated a javelin are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response;
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/note= "Javelin sequence"
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/note= "Javelin sequence"
Disclosure; Page 25; 150pp; English
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'note= "Linker"
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/note=
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(HOEM/) HOE M.
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Length 32,

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antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma antigen peptides of the invention
                                                                                                           Sequence 32
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Best Local S
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                                                                                                  number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The artigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is human gpl00 MHC class I mutant peptide antigen
                                                                              comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, 9p 100, MAGE antigens, BAGE antigens, NYEG1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma
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                                                                                   invention relates to an antigenic complex,
or prevent infectious disease or malignancy
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                                        Disclosure; Page 14; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU72245 standard; peptide; 32 AA.
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AL-AWQATI Q.
MAYHEW M.
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100.0%; Score 64; DB 4;
100.0%; Pred. No. 0:014;
ive 0; Mismatches (
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protein and a melanoma antigen i.e. tyrosinase.
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LIVINGSTON P.
AL-AWQATI Q.
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Best Local Similarity
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Matches

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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, OMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the the antigen should to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                                                                                                                                           Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYBS01; GN2; tyrosinase related protein 1; tyrosinase related protein 1; tyrosinase arelated protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                                                      gp100-derived melanoma antigen, javelin peptide #94.
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AAU72315 standard; peptide; 32
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                                                                               (first entry)
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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(HOEM/) HOE M.
                                                                                                                                                                                                                                                                       Homo sapiens.
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                                      AAU72315;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYE01, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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Indela
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Mismatches
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                                                                                                                                                                                   AAU72285 standard, peptide, 32 AA.
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Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
                                         HWDFAWPW 8
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
Synthetic.
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(ALAW/) (MAYH/) (HOEM/)

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Gaps

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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response;

RESULT 262 AAU72315

Best Loc Matches

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AAU72330 standard; peptide; 32 AA.
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                                                                                                                                                                                                                                                                                                                                                                                              cancer vaccine for the
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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LIVINGSTON P.
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Best Local Similarity
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immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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1es 8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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    Mayhew M,
                                                                                                                                                                                                                     protein and a melanoma antigen i.e. tyrosinase
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    Al-Awqati Q,
                                                                                                                                                                     treatment of
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Homo sapiens.
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                                                                                                                         The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase, related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                       Anti cancer vaccine for the treatment of melanoma comprises a heat shock protein and a melanoma antigen i.e. tyrosinase.
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                               Hoe M;
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                                Mayhew M,
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                               Noughton A, Livingston P, Al-Awgati Q,
                                                                                                        Disclosure; Page 27; 150pp; English
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                                                                                                                                                                                                                                     antigen peptides of the invention
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYES01, MART antigens, GMZ, antigence portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. ANUT1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                             The invention relates to a method of induction of an immune response,
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Disclosure; Page 29; 150pp; English
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antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAV71980-AAV72481 represent melanoma antigen peptides of the invention
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AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, BAGE antigens, NYESO1, MART antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAUT1980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                               AAU72440 standard; peptide; 32 AA.
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LIVINGSTON P.
AL-AWQATI Q.
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MAYHEW M.
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                                                                                                                                                                                                                                                                                                                                                                                                                     comportaing administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GMZ, antigenic portions and combinations of these. The melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AMVI1980-AAVI2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                              comprises a heat shock
                                                                                       Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherspetic; heat shock protein; tyrosinase; BAGE; NYSEG1; GNZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; davelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                     MAGE-1/3-derived melanoma antigen, javelin peptide #19.
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                                                                                                                                                                                                                                                                                                                                                                       protein and a melanoma antigen i.e. tyrosinase
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         AAU72460 standard; peptide; 32 AA
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LIVINGSTON P.
AL-AWQATI Q.
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AAU72460
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigens operations and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AUV11980-AAUV12481 represent melanoma antigen peptides of the invention
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Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunothezapeutic; heat shock protein; tyrosinase; BAGE; VRSB01; GWZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
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protein and a melanoma antigen i.e. tyrosinase.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, trainingent related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
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                                                                                Al-Awgati Q,
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(LIVI/) LIVINGSTON P.
(ALAW) AL-AWQATI Q.
(MAYH/) MAYHEW M.
(HOEM/) HOE M.
AL-AWQATI Q.
                                                                                                                         WPI; 2001-663092/76
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                   MAYHEW M.
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  ALAW/)
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protein and a melanoma antigen i.e. tyrosinase
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                                                           17-APR-2001; 2001WO-US012449.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Synthetic.
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                                                              invention relates to a method of induction of an immune response,
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protein and a melanoma antigen i.e. tyrosinase.
protein and a melanoma antigen i.e. tyrosinase
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                               Local Similarity
tes 8, Conserv
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                                                                                                                                                                                                                                                                 Sequence 32 AA;
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antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma AAU71980-AAU72481 represent melanoma antigen peptides of the invention
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
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Best Local Similarity
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                                                                                                                            Sequence 32 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer vaccine for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                              immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
                              Gaps
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100.0%; Score 64; DB 4; Length 32; 100.0%; Pred. No. 0.014;
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100.0%; Pred. No. 0.014;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                   gp100-derived melanoma antigen, javelin peptide #119.
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                             8; Conservative
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LIVINGSTON P.
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MAYHEW M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    melanoma comprises a heat shock
                                                                                                             Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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                                                                                     WAGE-1/3-derived melanoma antigen, javelin peptide #14.
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AAU72455 standard; peptide; 32 AA.
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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(HOUG/) HOUGHTON A.
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             Melanoma antigen, MART-1; MAGE-1; gp100; cytostatic; immune response; immuncherapetic; heat shock protein; tyrosinase; BAGE; NYBS01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T celle-1; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                    protein and a melanoma antigen i.e. tyrosinase
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, tyrosinase related protein 1, tyrosinase antigens (MZ, antigens, MZ, antigens, MZ, antigens of these: The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AMJ1980-AMJ12481 represent melanoma antigen peptides of the invention
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100.0%; Pred. No. 0.014;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                        Houghton A, Livingston P, Al-Awgati Q, Mayhew M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein and a melanoma antigen i.e. tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer vaccine for the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 31; 150pp; English
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17-APR-2001; 2001WO-US012449.
                                                                       17-APR-2000; 2000US-0197462P.
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                                                                                                                                      HOUGHTON A.
LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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Best Local Similarity
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comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 1, tyrosinase antigens, GM2, antigens, CM2, antigens, DA2, antigens, DA2, antigens, DA2, antigens and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANV1980-AANV2481 represent melanoma antigen peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of induction of an immune response, comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanome antigen, where the melanome antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase
  comprises a heat shock
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                                                                                                    invention relates to a method of induction of an immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MAGE-1/3-derived melanoma antigen, javelin peptide #24.
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Anti cancer vaccine for the treatment of melanoma protein and a melanoma antigen i.e. tyrosinase.
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protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 33; 150pp; English.
                                                            Disclosure; Page 28; 150pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
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(LIVI/)
(ALAW/)
(MAYH/)
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                                                                                                                                                                                                                                                                                                    comprising administration of an immunotherapeutic composition, comprising a heat shock protein, and a melanoma antigen, where the melanoma antigen is selected from tyrosinase, tyrosinase related protein 1, tyrosinase related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the heat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma. AAU71980-AAU72481 represent melanoma
                                                                                                                                                                                  for the treatment of melanoma comprises a heat shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunotherapeutic; heat shock protein; tyrosinase; BAGE; NYEs01; GM2; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; javelin molecule; melanoma antigen recognised by T cells-1; human.
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100.0%; Pred. No. 0.014;
Mismatches 0; Indels
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                                                                                                    Mayhew M,
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protein and a melanoma antigen i.e. tyrosinase.
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                                                                                                    Houghton A, Livingston P, Al-Awgati Q,
                                                                                                                                                                                                                                           Disclosure; Page 28; 150pp; English
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8; Conservative
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LIVINGSTON P.
AL-AWQATI Q.
MAYHEW M.
  LIVINGSTON P.
                     AL-AWQATI Q.
MAYHEW M.
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(LIVI/)
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RESULT 282

AAU72360

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Best Loc Matches

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Gaps

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100.0%; Score 64; DB 4; Length 32; 100.0%; Pred. No. 0.014; cive 0; Mismatches 0; Indels

8; Conservative

Query Match Best Local Similarity Matches 8; Conserv

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1 HWDFAWPW

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Chicken MHC class II peptide antigen #3.

12-FEB-2002 (first entry)

AAE13451;

AAE13451 standard; peptide; 38 AA.

AAE1345: RESULT

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related protein 2, gp 100, MAGE antigens, BAGE antigens, NYESO1, MART antigens, GM2, antigenic portions and combinations of these. The melanoma antigen is covalently bound to a javelin molecule, where the melanoma antigen bound to the javelin molecule is non-covalently bound to the hat shock protein. The composition is useful for inducing an immune response for the treatment of melanoma ANV11980-AAU72481 represent melanoma antigen peptides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Melanoma antigen; MART-1; MAGE-1; gp100; cytostatic; immune response; immunotherapatic; heat shock protain; tyrosinase; BAGE; NYSE01; GNZ; tyrosinase related protein 1; tyrosinase related protein 2; vaccine; Javelin molecule; melanoma antigen recognised by T cells-1; human.
                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MART-1-derived melanoma antigen, javelin peptide #24.
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                                                                                                                                                                                                                                                                                                                                                                                              AAU72385 standard; peptide; 32 AA.
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                                                                                                                                                                                                         Local Similarity 100.
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(LIVI/) LIVINGSTON P.
(ALAW/) AL-AWQATI Q.
(MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-663092/76
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                                                                                                                                                        Sequence 32 AA;
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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatability complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken MHC class II peptide antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                   Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatability complex; MHC; therapy; immune response; malignancy; chicken.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 5; Length 38;
                                                                                                                                                                                                                                                  II epitope"
                                                                                                                                                                                              'note= "Javelin sequence"
                                                                                                                                                                                                                                                                                      31. .38
/note= "Javelin sequence"
                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                 "MHC class
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HOE M.
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                                                                                                                                           Gallus gallus
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Sequence 32

us-10-053-520-143.rag

AAE13458 standard; protein; 100 AA

RESULT 287

AAE13458

AAE13458;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                                                                                                                                                                       Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatibility complex; MHC; therapy; immune response; malignancy; chicken.
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                         Indels
Pred. No. 0.016;
                                                                                                                                                                                                                                                                                                                                                                                      /note= "MHC class II epitope"
93. .100
                                                                                                                                                                                                                                                                                                                                                                  "MHC class I epitope"
                                                                                                                                                                                                                              Chicken ovalbumin derived protein domain #3.
                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Javelin sequence"
                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                              AAE13460 standard; protein; 100 AA.
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            100.08;
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                                                                                                                                                                                                   (first entry)
                         8; Conservative
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MAYHEW M.
HOE M.
                                                 HWDFAWPW 8
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HWDFAWPW 8
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           Similarity
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                                                                                                                                                                                                                                                                                                           gallus
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           Best Local
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(HOEM/)
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                                                                                                                                                                                                                                                                                                             Ballus
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                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                          Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatibility complex; MHC; therapy; immune response; malignancy; chicken.
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                                                                                                                                                                                                                                                                      'note= "MHC class II epitope"
                                                                                                                                                                                                                                          "MHC class I epitope"
                                                                    Chicken ovalbumin derived protein domain #1.
                                                                                                                                                                                                             /note= "Javelin sequence"
                                                                                                                                                                                 Location/Qualifiers
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                                        (first entry)
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/note= "N
74. .89
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                                                                                                                                                                                                                                                                                                                                                                                                                 (ROTH/) ROTHMAN J E. (MAYH/) MAYHEW M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-017594/02.
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Matches 8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                               HOE M.
                                                                                                                                                     Gallus gallus
                                        12-FEB-2002
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Best Loc Matches

AAE13461;

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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiological conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatibility complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken ovalbumin derived protein domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                             Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatibility complex; MHC; therapy; immune response; malignancy; chicken.
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                                                                                                                                                                                                                                                                                77. .92
/note= "MHC class II epitope"
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                                                                                                                                                                                                                                                              "MHC class I epitope"
Chicken ovalbumin derived protein domain #2
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'note= "Javelin Bequence"
                                                                                                                                                  location/Qualifiers
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                                                                                                                                                                                                                           "Linker
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                                                                                                                                                                                                                                                                                                                                                                                                                   17-APR-2001; 2001WO-US012567
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                                                                                                                                                                                                                                               69. .76
/note= "N
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/note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ROTH/) ROTHMAN J E. (MAYH/) MAYHEW M. (HOEM/) HOE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-017594/02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 103 AA;
                                                                                                                                                                                                                                                                                                                                           WO200179259-A1
                                                                                                              Gallus gallus.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
                                                                                                            Antigenic complex; epitope; heat shock protein; HSP; tether; javelin; major histocompatibility complex; MHC; therapy; immune response;
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                                                                                                                                                                                                                                                                                                    "MHC class II epitope"
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                                                                           Chicken ovalbumin derived protein domain #4.
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/note= "Javelin sequence"
                                                                                                                                                                                                                                                                                                                                           /note= "Linker sequence"
                                                                                                                                                                                                           Location/Qualifiers
58. .65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 14; 47pp; English.
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HWDFAWPW 103
                                                                                                                                                    malignancy; chicken.
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                                                                                                                                                                                       Gallus gallus
                                     12-FEB-2002
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(MAYH/) (HOEM/) (ROTH/)

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Query Match

Matches

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AAE13459 ID AAE1 XX AC AAE1 XX DT 12-F

RESULT 289

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                                                                                                                                                                                                                                                                                                                                                                                                              A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
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                                                                                                                                    "MHC class II epitope"
                                                                                                         "MHC class I epitope"
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                                                                            note= "Javelin sequence"
                                                                                                                                                             /note= "Javelin sequence"
                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 15; 47pp; English.
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                                                                                                                                     /note=
                                                                                                          /note=
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 malignancy; chicken
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                                                                                                                                                                                        40200179259-A1
                         Gallus gallus
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Gallus gallus

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The present invention relates to an antigenic complex, comprising a number of epitopes non-covalently joined to a heat shock protein (HSP) by a tethering molecule referred to as javelin which has affinity for the HSP under physiclogical conditions, where the epitopes are covalently joined to the tethering molecule and one epitope is major histocompatibility complex class I (MHC) and the other MHC class II. The antigenic complex is used to induce immune responses directed towards the treatment or prevention of infectious diseases and malignancies. The present sequence is chicken ovalbumin derived protein domain
                                                                                                                                                                                                                                                                                                                                                                                                    A new antigenic complex comprising epitopes non-covalently joined to a heat shock protein by a molecular tether designated a javelin are useful to treat or prevent infectious disease or malignancy.
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                                                                                                           /note= "MHc class II epitope"
                                                        104. .111
/note= "Javelin sequence"
                         /note= "Javelin sequence"
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cocation/Qualifiers
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                                                       /note= "Linker"
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Best Local Similarity
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(MAYH/) MAYHEW M.
(HOEM/) HOE M.
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OM protein - protein search, using sw model

Run on:

March 24, 2006, 07:44:44; Search time 39 Seconds (without alignments) 19.737 Million cell updates/sec

US-10-053-520-143 64 1 HWDFAWPW 8 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283416 segs, 96216763 residues Searched: Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES	Description	ATP-binding casset	Wnt-7a protein - m	hypothetical prote	probable sulfate t	mullerian inhibiti	SF16 protein, poll	hypothetical prote	transposase - Esch	carbamoy1-phosphat	endo-1,4-beta-xyla	hypothetical prote	A-alpha X protein		×	protein F47D12.9 [hypothetical prote	integral membrane	hypothetical prote	hypothetical prote	conserved hypothet	aldose epimerase f	hypothetical prote	malonyl-CoA decarb	conserved hypothet	cinnamoyl ester hy	probable membrane	probable two-compo	subtilisin-like pr	drug efflux transp
	OI E	A59189	2 G36470	2 B83262		: T11753			1 TQECT		2 T08469		2 S62748	S62749	S62750		2 AC1927			2 B85961	_		_		2 AG2872			: T36531	B3	E83002
	Query Match Length DB	1529		340					•	•	••	•	•••	537	538				217	•	•	•	309	• •					•	206
de	Query	68.8	67.2	9.59	65.6	65.6	64.1	64.1		62.5	62.5	62.5	62.5	62.5	62.5	62.5	61.7	6.09	6.09	6.09	6.09	6.09		6.09	60.9	6.09	6.09	6.09	6.09	60.9
	Score	44	43	42	42	42	41	41	41	40	40	40	40	40	40	4	39.8	39	39	39	39	39	39	39	39		39	39	39	39
		-	7	m	4	വ	9	7	80	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29

sodium/solute symp hypothetical prote	hypothetical prote probable formate d	probable transport probable transport hypothetical 34.5	hypothetical prote hypothetical prote		hypothetical prote NADH2 dehydrogenas	transcription fact hypothetical prote	photosystem II chl
A82432 G70941	T34482 A95262	G85876 A65008	E83128 G82611	T25957 T04056	T04055 B82822	T51834 T03996	S42647
0.0	000	9 79 79	01 02	0 0	01	00	7
513 515	521 1034	310	80 219	225	326 330	330	459
60.09	0.09	200	59.4	59.4	59.4 59.4	59.4	59.4
6 E	330	38.6	38	388	38	38	38
30 31	335	3 22	37	33 40	41 42	4 4 4	45

ALIGNMENTS

RESULT 1 A59189 ATP-binding cassette transporter - human (fragment) N;Alternate names: KIAA1062 protein C;Species: Homo sapiens (man) C;Date: 18-Feb-2000 #sequence_revision 18-Feb-2000 #text_change 31-Dec-2004	Circossion: Abylogase, T.; Ishikawa, K.; Hirosawa, M.; Miyajima, N.; Tanaka, A.; Kotani, P. R.; Kikuno, R.; Magase, T.; Ishikawa, K.; Hirosawa, M.; Miyajima, N.; Tanaka, A.; Kotani, P. DNA Res. 6, 197-205, 1999 A;Title: Prediction of the coding sequences of unidentified human genes. XIV. The complet A;Reference number: Z22861; MUID:99397452; PMID:10470851 A;Accession: A59119 A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-1529 KIKA A;Cross-references: UNIPROT:09HC28; UNIPARC:UDI000017A0E0; GB:AB028985; NID:g5689460; PII A;Experimental source: chromosome 9; clone hj03579; clone lib pBluescriptII SK plus; tiss	A; Worder: XIAA1062 A; Note: XIAA1062 A; Note: XIAA1062 Query Match Best Local Similarity 57.1%; Pred. No. 70; Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0; Qy 2 WDFAWPW 8 ::: Db 39 WEWSWPW 45		A;Reference number: A36470; MUID:91122634; PMID:2279700 A;Scaesaion: G36470 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-349 <gav> A;Cross-references: UNIPROT:P24383; UNIPARC:UPI0000029D72; GB:M89801; NID:g202409; PIDN:J C;Superfamily: int-1 transforming protein C;Superfamily: int-1 transforming protein Cuery Match Best Local Similarity 62.5%; Pred. No. 22; Best Local Similarity 62.5%; Pred. No. 22; Matches 5; Conservative 0; Mismatches 3; Indels 0; Gaps 0;</gav>
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1 HWDFAWPW 8

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N

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A; Molecule type: mRNA
A; Residues: 1-575 < DANS
A; Residues: 1-575 < DANS
C; Genetics:
A; Genetics:
A; Genetics:
A; Genetics:
A; Genetics:
A; Genetics:
C; Function:
Description: causes regression of the Mullerian duct, inhibits the growth of tumors desibition:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q39954; UNIPARC:UPI00000A035A; EMBL:X74772; NID:g560149; PIDN A;Bxperimental source: strain HA401B; pollen C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SF16 protein, pollen specific - common sunflower C; Species: Helianthus annuus (common sunflower) C; Species: Helianthus annuus (common sunflower) C; Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 09-Jul-2004 C; Accession: T13992 R; Dudareva, N.; Evrard, J.L.; Pillay, D.T.N.; Steinmetz, A. R; Dudareva, N.; Evrard, J.S.; Pillay, D.T.N.; Steinmetz, A. A; Title: Nucleotide Sequence of a pollen-specific cDNA from Helianthus annuus L encoding A; Reference number: Z17847; MUID:95062749; PMID:7972527
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C,Superfamily: inhibin
C,Keywords: cytotoxin; glycoprotein; gonadal differentiation; testis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 65.6%; Score 42; DB 2; Best Local Similarity 71.4%; Pred. No. 51; Matches 5; Conservative 1; Mismatches
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A;Status: preliminary; translated from GB/EMBL/DDBJ
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64.1%; Score 41; DB
Best Local Similarity 50.0%; Pred. No. 91;
Matches 4; Conservative 1; Mismatches
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A;Introns: 7/3; 239/1; 287/3; 398/3
C;Superfamily: COBRA-like protein 7 precursor
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247 HPDWAWSW 254
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47 HWDWNWP 53
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probable sulfate transporter - Chlamydia trachomatis
C;Species: Chlamydia trachomatis
C;Species: Unlamydia trachomatis
C;Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 09-Ju1-2004
C;Accession: A71463
Science 282, 754-759, 1998
A;Stephens, R.S.; Kalman, S.; Lammel, C.J.; Fan, J.; Marathe, R.; Aravind, L.; Mitchell,
Science 282, 754-759, 1998
A;Title: Genome sequence of an obligate intracellular pathogen of humans: Chlamydia trac
A;Reference number: A71570; MUID:99000809; PMID:9784136
A;Accession: A7140
                                                                                                                                                                                                                                                                                                                               C;Species: Pseudomonas aeruginosa
C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
C;Accession: B83262
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Bx adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathch; A;Reference number: A82950; MUID: 20437337; PMID: 10984043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
*Residues: 1-34 o.stro.
A;Cross-references: UNIPROT:Q9HZD5; UNIPARC:UPI00000C5900; GB:AE004731; GB:AE004091; NIC
A;Experimental source: strain PAO1
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                                                                                                                                                                                                                                                                                            hypothetical protein PA3073 [imported] - Pseudomonas aeruginosa (strain PAO1)
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mulication substitud factor precursor - pig
(;Species: Sus scrofa domestica (domestic pig)
(;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 09-Jul-2004
(;Accession: T1153
R;Daneau, 1:; Silversides, D.W.
aubmitted to the EMBL Data Library, July 1998
A;Reference number: Z17334
(Å;Accession: T11753
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         ||| | | |
HWDEGWKW 149
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174 YWDYLWTW 181
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les 5; Conserv
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2 FEFAWPW 8
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A,Gene: PA3073
                                                      142
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Matches
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Cispecies: Schizophyllum commune
Cipate: 24-Mag-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
Cipate: 24-Mag-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
Cipatesesion. S62748
R.Marion, A.L.; Bartholomew, K.A.; Wu, J.; Yang, H.; Novotny, C.P.; Ullrich, R.C.
Curr. Genet. 29, 143-149, 1996
A;Title: The A-alpha mating-type locus of Schizophyllum commune: structure and function c
A;Reference number: S62748 MUID:96418876; PMID:8821661
A;Reference number: S62748
A;Reference number: S62748
A;Residues: 1-532 - MAR>
A;Roberences: UNIPROT:002464; UNIPARC:UPI00006ACA9; EMBL:U13942; NID:g537620; PID)
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, August 1994
C;Genetics:
                                                                                                                                                                                            A;Cross-references: UNIPROT:Q12603; UNIPARC:UPI000013907F; EMBL:L39866; NID:g973982; PID:
A;Experimental source: strain Rt46B.1
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                          A; Description: hydrolyzes xylan to xylotriose and xylobiose but could not hydrolyze xylot (S.Guperfamily: xylanase; Streptomyces ando-1,4-beta-xylanase. A homology C; Keywords: glycosidase; hydrolase; polysacoharide degradation F; 60-352/Domain: Streptomyces endo-1,4-beta-xylanase A homology <SXX>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,Cross-references: UNIPROT:004629; UNIPARC:UPI00000A2ABB; EMBL:AF007270; NID:G2191157; N
Appl. Environ. Microbiol. 61, 4403-4408, 1995
Affile: Clohing, sequencing, and expression of a xylanase gene from the extreme thermopl
A;Reference number: Z16432; MUID:96086022; PMID:8534104
A;Accession: T08469
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C;Species: Arabidopsis_thaliana (mouse-ear cress)
C;Pate: 19-Feb-1999 #sequence_revision 19-Feb-1999 #text_change 09-Jul-2004
C;Accession: T01762
R;Miller, N.; Beck, C.; Kramer, J.
submitted to the EMBL Data Library, June 1997
A;Description: The sequence of A. thaliana IG002P16.
A;Reference number: 214421
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                                                                                                                 A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-352 <GIB>
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Best Local Similarity 83.3%; Pred. No. 85;
Matches 5; Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 40; DB
Pred. No. 61;
0; Mismatches
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A,Status: translated from GB/EMBL/DDBJ
A,Rolecule type: DNA
A,Residues: 1-482 <MIL>
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Best Local Similarity 71.4%;
Matches 5; Conservative (
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A,Gene: ATSP:A IG002P16.6
A,Map position: 5
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R;Heffron, F.; McCarthy, B.J.; Ohtsubo, H.; Ohtsubo, B.
R:Heffron, F.; McCarthy, B.J.; Ohtsubo, H.; Ohtsubo, B.
A;Title: DNA sequence analysis of the transposon Tn3: three genes and three sites involv A;Reference number: A90784; MUID:80090058; PMID:391406
A;Accession: A03538
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A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              carbamoyl-phosphate synthase (glutamine-hydrolysing) (EC 6.3.5.5) [imported] - Brucella C;Species: Brucella melitensis
C;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:Q8YEUG; UNIPARC:UPI0000581D9; GB:AE008917; PIDN:AAL52962.1;
A;Experimental source: strain 16M
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C;Species: DictyOglomus thermophilum
C;Species: DictyOglomus thermophilum
C;Accession: T0469
R;Gibbs, M.D.; Reeves, R.A.; Bergquist, P.L.
                                                                                                                                                                                                                                  transposase - Escherichia coli transposon Tn3
C;Species: Escherichia coli
C;Date: 30-Jun-1980 #sequence_revision 15-Oct-1982 #text_change 09-Jul-2004
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A;Residues: 1-1015 <HEF>
A;Cross-references: UNIPROT:P03008; UNIPARC:UPI00001370D7
C;Comment: This protein is required for transposition of transposon Tn3.
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Pred. No. 1.3e+02;
1; Mismatches 0;
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C, Keywords: DNA binding; DNA replication
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Best Local Similarity 71.4%;
Matches 5; Conservative
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Best Local Similarity 83.39
Matches 5; Conservative
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EFAWPW 133
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FSFAWPW 51
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A;Molecule type: DNA
A;Residues: 1-237 <KUR>
                               1 HWDFAWPW
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C; Keywords: ligase
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C; Superfamily: A-alpha X protein

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A; Map position: 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues: 1-538 <MAR>
A;Cross-references: UNIPROT:Q02467; UNIPARC:UPI000069D65; EMBL:U13944; NID:g537624; PID A;Note: the nucleotide sequence was submitted to the EMBL Data Library, August 1994 C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                  RiMarion, A.L.; Bartholomew, K.A.; Wu, J.; Yang, H.; Novotny, C.P.; Ullrich, R.C. Cur. Genet. 29, 143-149, 1996
A;Itle: The A-alpha mating-type locus of Schizophyllum commune: structure and function A;Reference number: S62748; MUID:96418876; PMID:8821661
A;Accession: S62749
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A;Title: The A-alpha mating-type locus of Schizophyllum commune: structure and function A;Reference number: S62748; MUID:96418876; PMID:8821661
A;Accession: S62750
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
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                                                                                                                                                                                                                                                                                               A-alpha X protein (allele 3) - bracket fungus (Schizophyllum commune)
C;Species: Schizophyllum commune
C;Date: 24-Aug-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S62749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Schizophyllum commune
C;Date: 24-Aug-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S62750
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'Match 62.5%; Score 40; DB 2; Length 532; Local Similarity 83.3%; Pred. No. 94; local Si Conservative 0; Mismatches 1; Indels
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C;Superfamily: A-alpha X protein
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C,Superfamily: A-alpha X protein
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les 5; Conservative
                                                                                                                                                                   197 WDFQWP 202
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     Query Match
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C)Accession: B88479
R;anonymous, The C. elegans Sequencing Consortium.
Science 282, 2012-2018, 1998
A;Title: Genome sequence of the nematode C. elegans: a platform for investigating biology
A;Fitle: Genome sequence of the nematode C. elegans: a platform for investigating biology
A;Reference number: A75000, MUID:99069613; PMID:8851916
A;Note: see websites genome.wustl.edu/gsc/C_elegans/ and www_sanger.ac.uk/Projects/C_elegans/ A;Note: published errata appeared in Science 283, 35, 1999; Science 283, 2103, 1999; and
A;Accession: B88479
A;Accession: brainmary
A;Molecule type: DNA
A;Residues: 1-545 <STO>
                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:Q09392; UNIPARC:UP1000013BF71; GB:chr_III; PIDN:AAA64320.1; I
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Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004
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Pred. No. 96;
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Job time : 42 secs
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05w368 oryza sativ
05w366 oryza sativ
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05w366 oryza sativ
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                                                               March 24, 2006, 07:41:01; Search time 231 Seconds (without alignments) 24.434 Million cell updates/sec
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Q4kd40
Q5gtx5
Q8psf8
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Q82B30 STRAW
Q4UU5-XANCP
Q8D944 XANCP
Q85B1 GRYSA
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Larbig K.D., Christmann A., Johann A., Klockgether J., Hartsch T.,
Merkl R., Wiehlmann L., Fritz H.J., Tummler B.;
"Gene islands integrated into tRNA(Gly) genes confer genome diversity
on a Pseudomonas aeruginosa clone.";
J. Bacteriol. 184:665-6680(2002).
EMBL; AR40523; AAN62139.1; -; Genomic DNA.
GO; GO:0030288; C:periplasmic space (Sensu Gram-negative Bact. . .; IEA.
GO; GO:0015035; F:protein disulfide oxidoreductase activity; IEA.
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PubMed=15980861; DOI=10.1038/nbt1110;
PubMed=15980861; DOI=10.1038/nbt1110;
Paulsen I.T. Press C., Ravel J., Kobayashi D., Myers G.S.,
Mavrodi D., DeBoy R.T., Seshadri R., Ren Q., Madupu R., Dodson R.J.,
Durkin S., Brinkac L.M., Daugherty S.C., Sullivan S.A., Roscovitz M.,
Gwinn M.L., Zhou L., Nelson W.C., Weidman J., Watkins K., Tran K.,
Khouri H.M., Plerson E., Plerson L. III, Thomashow L., Loper J.,
"Complete genome sequence of the plant commensal Pseudomonas
fluorescens PE-5.";
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Q4KKC3_PSEF5 PRELIMINARY; PRT; 419 AA.
Q4KKC3_PSEF5 PRELIMINARY; PRT; 419 AA.
Q4KKC3_BSEP-2005 (TEMBLrel. 31, Last sequence update)
13-SEP-2005 (TEMBLrel. 31, Last annotation update)
13-SEP-2005 (TEMBLrel. 31, Last annotation update)
Intercellular adhesion protein A.
Indexcellular adhesion protein A.
Pseudomonas fluorescene (strain Pf-5).
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               70.3%; Score 45; DB 2; Length 254; 71.4%; Pred. No. 96; ive 0; Mismatches 2; Indels
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EMBL; CP000076; AaY95575.1; -; Genomic DNA.
SEQUENCE 419 AA; 48273 MW; C1784CCEOAF1B252 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     254 AA; 27649 MW; 212C116B689B8069 CRC64;
                                                                                                                                        01-MAR-2003 (TrEMBLrel. 23, Created)
1-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Hypothetical protein ORF C46.
                                                    254 AA
                                                    PRT;
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Best Local Similarity 71.4%;
Matches 5; Conservative
                                    QBGQ64 PSEAE PRELIMINARY,
QBGQ64;
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PSEAE
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RA Makamura Y., Kaneko T., Sato S., Ikeuchi M., Katoh H., Sasamoto S., Ratanabe A., Iriguchi M., Kawashima K., Kimura T., Kishida Y., Shimpo S., Kohara M., Matsumoto M., Matsumo A., Nakzaki N., Shimpo S., Sugimoto M., Takeuchi C., Yamada M., Tabata S.; Thermophilo S., Sugimoto M., Takeuchi C., Yamada M., Tabata S.; Thermophilo C., Yanada M., Tabata S.; Thermosynachococcus elongatus BP-1."; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Takamosynachococcus elongatus BP-1."; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Tabata S.; Danaga M., Managa M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arthrobacter sp. FB24.
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
Micrococcineae; Micrococcaceae; Arthrobacter.
NCBI_TaxID=290399;
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EMBL; AAHG01000024; EAL94587.1; -; Genomic DNA.
SEQUENCE 301 AA; 32541 MW; DD0C8E930AS7942E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Complete proteome; Transmembrane; Transport.
SEQUENCE 283 AA; 31039 MW; A3C74E30224EC4C7 CRC64;
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Pred. No. 1.5e+02;
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US DOE Joint Genome Institute (PGF-ORNL);
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TIGRPAMS; TIGR00969; 3a0106802; 1.
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62.5%;
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Q4NB48;
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STRAIN=FB24;
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Q4NB48
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         -i- CATALYTIC ACTIVITY: Obtusifoliol + 3 O(2) + 3 NADPH = 4-alpha-methyl-5-alpha-ergosta-8,14,24(28)-trien-3-beta-ol + formate + 3 NADP(+) + 3 H(2)).

-i- SIMILARITY: Balongs to the cytochrome P450 family.

EMBL; AACU01000440; EAA54287.1; -; Genomic_DNA.

InterPro; IPR001128; Cytochrome_P450.

InterPro; IPR00131; EP4501V.

PRINTS; PR00465; EP4501V.

PR0046501V.

PR0046501V.

PR0046501V.

PR0046501V.

PR0046501V.

PR00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Arch Midrobiol. 183:27-36(2005).
EMBL; CRS55306; CAI09711.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=EbN1;
PubMed=15551059; DOI=10.1007/s00203-004-0742-9;
Rabus R., Kube M., Heider J., Beck A., Heitmann K., Widdel F., Reinhardt R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Proteobacteria; Betaproteobacteria; Rhodocyclales;
Rhodocyclaceae; Azoarcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=nosR; OrderedLocusNames=AZOSEA35860; ORFNames=ebB224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 71.4%; Pred. No. 79; 5; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         2; Length 526;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synechococcus elongatus (Thermosynechococcus elongatus).
Bacteria, Cyanobacteria, Chroococcales, Synechococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     142 AA; 15770 MW; 8733EBB6E312A50D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                         70.3%; Score 45; DB 2; Li
100.0%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            142 AA
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Name=cysT, OrderedLocusNames=tlr0275;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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MEDLINE=22225144; PubMed=12240834;
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 70.3%; Best Local Similarity 100.0%; Matches 6; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-FEB-2005 (TrEMBLrel. 29, 01-FEB-2005 (TrEMBLrel. 29, 01-FEB-2005 (TrEMBLrel. 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Azoarcus sp. (strain EbN1)
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QBDM48;
01-MAR-2003 (TrEMBLrel. 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QSNZ03 AZOSE PRELIMINARY;
QSNZ03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 484 WDFAWP 489
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SEQUENCE 142 AA;
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Matches 5; Conserv
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QSW9GS_HUM
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Matches
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                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oryza sativa (japonica cultivar-group).
Sukaryota, Viridiplantes, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnollophyta, Liliopsida, Poales, Poaceae,
Ehrhartoideae, Oryzeae, Oryza.
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Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
Ehrhartoideae, Oryzeae, Oryza.
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"In-depth view of structure, activity, and evolution of rice chromosome 10.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           h 68.8%; Score 44; DB 2; Length 323; Similarity 71.4%; Pred. No. 1.7e+02; 5; Conservative 1; Mismatches 1; Indels
                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AC087182; AAL59037.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gramene, Q7XD95; -.
Hypothetical protein.
SEQUENCE 323 AA; 33440 MW; 5D63175DA9CD17D8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                               01-OCT-2003 (TrEMBLrel. 25, Created)
01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
10-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein.
ORFNames-OSJNBa0029C15.22;
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein OSJNBa0029C15.22.
      Pred. No. 1.6e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                         PRT;
71.48;
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QBW368 ORYSA PRELIMINARY;
Q8W368;
                                                                                                                                                                                                                                                                                                                                  Q7xd95 ORYSA PRELIMINARY;
Q7xd95;
Best Local Similarity 71.4 Matches 5; Conservative
                                                                                                                                                                       240 WDATWPW 246
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69 WPFSWPW 75
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                                                                                                   2 WDFAWPW 8
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Best Local Similarity
Matches 5; Conserv
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Homo saptens (Human).
Eukaryota, Merazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                          Gaps
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                                                                         Length 323;
                                                                                                          Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NON TER 1 1
SEQUENCE 1440 AA; 159853 MW; 27F20F0D7E87B8FD CRC64;
Gramene, Q8W368; -.
Hypothetical protein.
SRQUENCE 323 AA; 33440 MW; 5D63175DA9CD17D8 CRC64;
                                                                                                                                                                                                                                                               Q5W9G5, HUMAN PRELIMINARY; PRT; 1440 AA.
Q5W9G5,
01-FEB-2005 (TrEMBLrel. 29, Created)
01-FEB-2005 (TrEMBLrel. 29, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
KIAA1062 splice variant 1 (Fragment).
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QTGMWT;
QTGMWT;
QTGMWT;
OS-JUL-2004 (TrEMBLrel. 27, Last sequence update)
OS-JUL-2004 (TrEMBLrel. 27, Last sequence update)
KIAA1062 protein (Fragment).
                                                                    / Match 68.8%; Score 44; DB 2; Le Local Similarity 71.4%; Pred. No. 1.7e+02; nes 5; Conservative 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00211; ABC_TRANSPORTER_1; UNKNOWN_1.
PROSITE; PS50893; ABC_TRANSPORTER_2; 1.
PROSITE; PS00022; BGF_1; UNKNOWN_1.
ACCORDER; PS00213; LIPOCALIN; UNKNOWN_1.
ATP-binding; Nucleotide-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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295 WEWSWPW 301
                                                                                                                                                                | |:|||
69 WFFSWPW 75
                                                                                                                                             2 WDFAWPW 8
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Transport
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CARBOHYD
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TISSUE=Brain;
MRDLINFS=20427713; PubMed=10970803; DOI=10.1042/0264-6021:3500865;
Zhao L.-X., Zhou C.-J., Tanaka A., Nakata M., Hirabayashi T.,
Amachi T., Shioda S., Ueda K., Inagaki N., Hirabayashi T.,
Amachi T., Shioda S., Ueda K., Inagaki N., Hirabayashi T.,
Piloning, characterization and tissue distribution of the rat ATP-
binding cassette (ABC) transporter ABC2/ABCA2.";
Biochem. J. 350:865-872 (2000).
-!- FUNCTION: Probable transporter, its natural substrate has not been
found yet. May have a role in macrophage lipid metabolism and
neural development.
-!- SUBCELLULAR LOCATION: Integral membrane protein (Potential).
           Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                               TISSUE=Brain;
MEDLINE=99397452; PubMed=10470851;
Kikuno R., Nagase T., Ishikawa K.-I., Hirosawa M., Miyajima N.,
Tanaka A., Kotani H., Nomura N., Ohara O.;
"Prediction of the coding sequences of unidentified human genes. XIV.
The complete sequences of 100 new cDNA clones from brain which code
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rattus norvegicus (Rat).
Bukaryota, Metazoa, Chordata, Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-OCT-2004 (Rel. 45, Created)
25-OCT-2004 (Rel. 45, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
ATP-binding casette sub-family A member 2 (ATP-binding cassette transporter 2) (ATP-binding cassette Name-Abca2; Synonyms-Abc2;
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                                                                                                                                                                                                                                                                                                                                                                                                                             68.8%; Score 44; DB 2; Length 1771;
                                                                                                                                                  for large procedure in vitro.";

DNA Res. 6:197-205(1999).

EMBL; AB028985; Babagou to the ABC transporter family.

EMBL; AB028985; BAA63014.2; "RNA.

ENSEMD!; ENSG0000107331; Homo sapiens.

GO; GO:0005524; F:ATP binding; IEA.

GO; GO:0016887; F:ATPsase activity; IEA.

GO; GO:0000166; F:nucleotide binding; IEA.

InterPro; IPR003593; AAA ATPsase.

InterPro; IPR004299; AGF like.

InterPro; IPR002345; Lipocalin.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 7.8e+02;
3; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT; 2434 AA.
                                                                                                                                                                                                                                                                                           Pfan, PF00005; ABC tran; 2.
ProDom; PD000006; ABC transporter; 2.
SWART; SM00382; AAA; 2.
SWART; SM00311; ABC TRANSPORTER 1; 1.
PROSITE; PS50893; ABC TRANSPORTER 2; 2.
PROSITE; PS508021; BGF 1; UNKNOWN 1.
PROSITE; PS000213; LIPOCALIN; UNKNOWN 1.
ATP-binding; Nucleotide-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                          57.1%;
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Homo sapiens (Human)
                                                                     NUCLEOTIDE SEQUENCE
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                                              NCBI_TaxID=9606;
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ABCA2_RAT
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
TISSUE SPECIFICITY: Expressed at high levels in brain, at moderate levels in heart, kidney and lung, and at low levels in skeletal muscle, stomach, spleen, colon and pancreas. Not detected in the liver or small intestine. In brain, highly expressed in white matter and detected in oligodendrocytes.
                                                                                                                                                                                                                                                                                                                       GO; GO:0043190; C:ATP-binding cassette (ABC) transporter complex; ISS. GO; GO:0016021; C:integral to membrane; NAS. GO; GO:0005524; F:ATP binding; IDA. GO; GO:0005529; F:ATP binding; IDA. GO; GO:0006629; P:lipid merabolism; NAS. GO:0006810; P:transport; NAS.
                                                                                         -1- PTM: N-glycosylated.
-1- SIMILARITY: Belongs to the ABC transporter family, ABCA subfamily
-1- SIMILARITY: Contains 2 ABC transporter domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATP-binding; Glycoprotein; Nucleotide-binding; Repeat; Transmembrane;
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ABC transporter 2.
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ATP 2 (By similarit
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Ensembl; ENSRNOG0000014268; Rattus norvegicus.
RGD; 620238; Abca2.
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InterPro; IPR003439; AAA ATPase.
Pfam; PF00005; ABC tran; 2.
ProDom; PD000006; ABC transporter; 2.
PROSTTS; PS00211; ABC TRANSPORTER 1; 1.
PROSITE; PS50893; ABC TRANSPORTER 1; 1.
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AAK14335.1;
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                        Gaps
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Q9B2C7;

Q16-OCT-2001 (Rel. 40, Created)

16-OCT-2001 (Rel. 48, Last amnotation update)

13-SEP-2005 (Rel. 48, Last annotation update)

ATP-binding cassette sub-family A member 2 (ATP-binding cassette transporter 2) (ATP-binding cassette 2).

Name-ABCAZ; Synonyms-ABCZ;

Homo sapiens (Human).
1776 1776 N-linked (GlCNAC. . .) (Potential) 2055 2055 N-linked (GlCNAC. . .) (Potential) 2434 AA, 270928 MW; CD424A9C4F63513F CRC64;
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Pred. No. 1e+03;
3; Mismatches 0; Indels
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Pred. No. 1e+03;
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PROSITE; PS50893; ABC TRANSPORTER 2; 2.
PROSITE; PS00022; EGF 1; UNKNOWN 1.
PROSITE; PS00213; LIPÖCALIN; UNKÖNWN 1.
ATP-binding; Nacleotide-binding; Repeat; Transmembrane.
SEQUENCE 2435 AA; 269873 MW; 0C4ABE9087BECSC4 CRC64;
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Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databases.
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57.1%; Pred. No. 10...
3; Mismatches
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ProDom; PD000006; ABC transporter; 2.
SMART; SM00382; AAA; 2.
                                                                                                 h 68.8%;
Similarity 57.1%;
4; Conservative
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945 WEWSWPW 951
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NCBI_TaxID=9606;
  CARBOHYD
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Q5SPY5 HUMAN
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatical Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                      of
in
                                                                         NUCLEOTIDE SEQUENCE.
PubMed=11178988; DOI=10.1006/Dbrc.2001.4305;
PubMed=11178988; DOI=10.1006/Dbrc.2001.4305;
Maninski W.E., Piehler A., Pullmann K., Porsch-Oezcueruemez M., Duong C., Bared G.M., Buchler C., Schmitz G., "Complete coding sequence, promoter region, and genomic structure the human ABCA2 gene and evidence for sterol-dependent regulation
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EMBL, AF327659; AAK14335.1; OINED; Genomic_DNA.
EMBL, AF377669; AAK14335.1; JOINED; Genomic_DNA.
EMBL, AF327661; AAK1435.1; JOINED; Genomic_DNA.
EMBL, AF327661; AAK1435.1; JOINED; Genomic_DNA.
EMBL, AF327663; AAK14315.1; JOINED; Genomic_DNA.
EMBL; AF327663; AAK14335.1; JOINED; Genomic_DNA.
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PRT; 2436 AA

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Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
          Q9HC28;
01-MMR-2001 (TrEMBLrel. 16, Created)
01-MMR-2001 (TrEMBLrel. 16, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
ATP-binding cassette sub-family A member 2 (ABC transporter
 Q9HC28_HUMAN PRELIMINARY;
                                                                                                                                  NUCLEOTIDE SEQUENCE.
                                                                                                              WCBI_TaxID=9606;
                                                                                                                                            rissum=Brain;
                                                                  Name=ABCA2;
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                                                         ABCA2)
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                                   GO; GO:0043190; C:ATP-binding cassette (ABC) transporter complex; NAS.
GO; GO:0016021; C:integral to membrane; NAS.
GO; GO:0006621; C:integral to membrane; NAS.
GO; GO:0006610; P:transport; NAS.
InterPro; IPR003593; AAA ATPase.
InterPro; IPR003439; ABC transporter;
Pfam; PF00005; ABC trans; 2.
Probom; P0000066; ABC transporter; 2.
SWART; SM00382; AAA; 2.
                                                                                                                     PROSITE; PS00211; ABC_TRANSPORTER 1; 1.
PROSITE; PS50893; ABC_TRANSPORTER 2; 2.
ATP-binding; Glycoprotein; Nucleotide-binding; Repeat; Transmembrane;
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Pred. No. 1e+03;
3; Mismatches 0; Indels
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ABC transporter 2.
ATP 1 (Potential).
ATP 2 (Potential).
N-linked (GlCMAC.)
EMBL; AF327704; AAK14335.1; JOINED; Genomic_DNA.
Ensembl; ENSG0000107331; Homo sapiens.
HGNC; HGNC:32; ABCA2.
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Elochem. Biophys. Res. Commun. 281:249-258 (2001).

Elochem. Biophys. Res. Commun. 281:249-258 (2001).

Elochem. Biophys. Res. Commun. 281:249-258 (2001).

C. -1. SUBCELLULAR LOCATION: Integral membrane protein (By similarity).

EMBL; AF178941; AAG09372.1; -; mRNA.

EMBL; AF178941; AAG09372.1; -; mRNA.

REMBL; AF18941; AAG09372.1; -; mRNA.

REMBL; AF178941; AAG09372.1; -; mRNA.

REMBL; FIPTOROSS24; FATPEASE activity; IEA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0016021; R:ATPEASE activity; IEA.

RO; GO:000166; F:nucleotide binding; IEA.

RINTERPO: IPR003393; AAA, ATPEASE.

RINTERPO: IPR003495; AAA, ATPEASE.

RINTERPO: IPR00006; ABC transporter; 2.

EMBRT; SMO0382; AAA; Z.

EMBRT; SMO0382; AAA; Z.

EMBL; SMART; SMO0382; AAA; Z.
                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
MEDLINE=21092814; PubMed=11178988; DOI=10.1006/bbrc.2001.4305;
Kaminski W.E., Piehler A., Pullmann K., Porsch-Oezcueruemez M.,
Duong C., Bared G.M., Buchler C., Schmitz G.;
"Complete coding sequence, promoter region, and genomic structure of
the human ABCA2 gene and evidence for sterol-dependent regulation in
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MEDLINE=21205831; PubMed=11309290;
Vulevic B., Chen Z., Boyd J.T., Davis W. Jr., Walsh E.S.,
Belinsky M.G., Tew K.D.;
"Cloning and characterization of human adenosine 5'-triphosphate-
binding cassette, sub-family A, transporter 2 (ABCA2).";
Cancer Res. 61:3339-3347(2001).
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PROSITE; PS50893; ABC_TRANSPORTER_2; 2.
PROSITE; PS00022; EGF 1; UNKNOWN T.
PROSITE; PS00213; LIPGCALIN; UNKNOWN 1.
ATP-binding; Nucleotide-binding; Repeat; Transmembrane.
SEQUENCE 2436 AA; 269958 WW; E044A3AF14EA25D1 CRC64;
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Job time : 235 secs
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; Sequence 1, Application US/09794529B
; Patent No. 6641812
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.

                                                                                                          March 24, 2006, 12:32:02 ; Search time 46 Seconds (without alignments) 14.378 Million cell updates/sec
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Biocceleration Ltd.
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/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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US-09-794-517A-1

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US-09-52-168-1

US-09-536-295-1

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US-09-794-517A-9

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US-09-680-806A-9

US-09-794-629B-10

US-09-52-868-9

US-09-536-295-9

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                 version :
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                                                                                 protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                 GenCore
(c) 1993
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Maximum Match 100%
Listing first 1000
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Maximum DB seq length: 200000000
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Query
Match
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AAQQU Sequence Seq Sequence US-09-794-529B-16
US-09-794-529B-17
US-09-794-529B-19
US-09-794-517A-10
US-09-794-517A-11
US-09-794-517A-11
US-09-794-517A-11
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US-09-794-517A-15
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US-09-794-612-11
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US-09-680-680-680-11
US-09-552-680-11
US-09-636-295-11
US-09-636-295-11

ALIGNMENTS

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GENERAL INFORMATION:
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            APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                 - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: heat shock binding sequence; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 64; DB 2; I
ilarity 100.0%; Pred. No. 4.6e+05;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERÍCCT
CURRENT APPLICATION DATA:
FILLING DATE: 09-Jan-2002
CLASSIFICATION: 336
PRIOR APPLICATION: DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
FILING DATE: 40002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: FEbruary 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 425-5288
TELEX: «UNKNOWN)
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: cunknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                     COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWDFAWPW 8
GENERAL INFORMATION:
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Matches 8; Conserv
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RESULT 2 0S-09-14-517A-1 'Sequence 1, Application US/09794517A '; Patent No. 6656679

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APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                 - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; SEQUENCE DESCRIPTION: heat shock binding sequence; SEQUENCE DESCRIPTION; SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM COMPABLIA

COMPUTER: IBM COMPABLIA

SOFTWARE: Word Perfect

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/794,517A

FILING DATE: 19-OCC-2001

CLASSIFICATION DATA:

APPLICATION NUMBER: 60/002,479

FILING DATE: ANGUET 18, 1995

APPLICATION NUMBER: 60/002,490

FILING DATE: ANGUET 18, 1995

APPLICATION NUMBER: 00/001,645

FILING DATE: ANGUET 16, 1996

APPLICATION NUMBER: 09/011,645

FILING DATE: FEBRUARY 13, 1998

ATTORNEY/AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: 28,839

REFERENCE/DOCKET NUMBER: 28,839

REFERENCE/DOCKET NUMBER: 28,839
                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-011-645E-1; Sequence 1, Application US/09011645E; Patent No. 6663868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 425-720(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORWATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                   COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                         STREET: One Broadway
                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                          CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 HWDFAWPW 8
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Gaps
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                                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                  - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 4.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; SEQUENCE DESCRIPTION: heat shock binding sequence; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Diskette - 3.5 inch, 1
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKGT NUMBER: 11746/1
                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAXECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/09680806A Patent No. 6719974 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: mouse
                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-680-806A-1
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                                                                                                                                                                                                                                                                                                             STATE: NO.
STATE: NO.
STATE: NO.
ZOUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13.Feb-1998
CLASSIFICATION ATA:
PRIOR APPLICATION DATA:
                                       ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                        APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/09794832
Sequence 1. Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 4.6e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Deflucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-011-645E-1
                                                                                                                                                                                                        NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
INPORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS
LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: mouse
                                                                                                                                                                                                                                                                                              CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 HWDFAWPW 8
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  GENERAL INFORMATION:
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Matches 8; Conserv
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TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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; Sequence 1, Application US/09636295
; Patent No. 6773707
; GENERAL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 4.6e+05; iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: heat shock binding sequence SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: DBRECTE - 3.5 INCH, 1.4
COMPUTER: DBM COMPABLIDE
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                          NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: cunknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: mouse
                                                                                                                                                                                                                                                                    CITY: New York
                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 HWDFAWPW 8
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ROTHMAN, James E.
HARTL, P. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark Shock Protein-Based Vaccines and
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'Sequence 1, Application US/09552868

'Patent No. 6761892

GENERAL INFORMATION:

APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                ZIP: 10004

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:

FILING DATE: 05-0ct 2000

CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 4.6e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: heat shock binding sequence SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
                                                                                                                                                          Immunotherapies
                                                                                                                                                                          NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: <UNKNOWN >
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                               STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: mouse
                                                                                                                                                                                                                                                                  CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |||||||||||||
HWDFAWPW 8
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TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
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Patent No. 6641812
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 2; Length 19; 100.0%; Pred. No. 0.0027; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: MS DOS
SOFWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-794-529B-8
                                                                         HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 425-5288
           ROTHMAN, James E
HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                 ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                               STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: <Unknown>
                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
                                   HARTL, F. U
HOE, Mee H.
                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8; Conservative
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -09-794-529B-9
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ROTHWAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKCHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/09794529B
Sequence 8, Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                     STATE: NY
COMPUTER: 100005
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 64; DB 2; Length 8; 100.0%; Pred. No. 4.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: heat shock binding sequence; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
RILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATTON NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FELEFAX: (212) 425-5288
                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                   NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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; Sequence 9, Application US/09794517A
; Patent No. 6656679
; GENERAL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                     ZIF: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 100.0%; Score 64; DB 2; Length 19; Similarity 100.0%; Pred. No. 0.0027; 8; Conservative 0; Mismatches 0: Indele
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM COMPACTURE STATEM, MS DOS SOFTWARE: WORD PERFOR SOFTWARE: WORD PERFOR CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/794,517A FILING DATE: 19-Oct-2001
CLASSIFICATION S36
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/002,479
FILING DATE: ANGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: ANGUSE 18, 1995
APPLICATION NUMBER: PCT/USS6/13363
FILING DATE: ANGUSE 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDFUGATY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       )
SEQUENCE DESCRIPTION: SEQ 1D NO: 8:
US-09-794-517A-8
                                                                                                                                                                               NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS;
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, YOSHIZUMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDENESS: «Unknown»
TOPOLOGY: linear
WOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NOWS SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: no
PRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: Diskette - 3.5 inch, 1.44 Mb storage
SOFTWARE: WAS Description of Proceedings of Prizer Application NUMBER: 08/09/794,5298
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: AUGUST 16, 1996
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: PEDATUST 11, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                   TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 10
US-09-794-517A-8
; Sequence 8, Application US/09794517A
; Patent No. 6656679
; GENERAL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match

100.0%; Score 64; DB 2; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.0027;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) SEQUENCE DESCRIPTION: ) SEQ ID NO: 9: US-09-794-529B-9
                                                                                                                                                                                                                          ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                               NUMBER OF SEQUENCES: 3(CORRESPONDENCE ADDRESS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IYPE: amino acid
                                                                                                                                                                                                                                                                      CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                  COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HWDFAWPW 8
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Protein-Based Vaccines and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 64; DB 2; Length 19; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                        3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/011,645E FILING DATE: 13-Feb-1998 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                         HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Prote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-011-645E-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            425-5288
                   HARTL, F. Ulrich
HOE, Mee H.
                                                                                                                                                                                                                            STREET: One Broadway
CITY: New York
                                                                                                                                                                NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-011-645E-9
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                   HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKEHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5-09-011-645E-8
Sequence 8, Application US/09011645E
Patent No. 6661868
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                        - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 100.0%; Score 64; DB 2; Length 19; Similarity 100.0%; Pred. No. 0.0027; 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4

COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILLING DATE: 19-OCt-2001
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/U396/13363
FILING DATE: August 16, 1996
FILING DATE: PEDFUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: DeLocia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-5288
TELEFAX: (212) 425-5288
                                                                                                                                                 Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO: 9: US-09-794-517A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
ROTHMAN, James E
HARTL, F. Ulrich
                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION:
                                                                                                                                                                    SEQUENCES:
                                                                                                                                                                                                                                               CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                            COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                  NUMBER OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE
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COMPUTRY: US
COMPUTRY: US
COMPUTRY: US
COMPUTRY: 10004
COMPUTRY: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794.832
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown>
PRIOR APPLICATION: «Unkner: US/99/011,645
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGRYT INFORMATION:
NAME: DeLUCIA, RICHARD I.
REGISTRATION NUMBER: 28, 839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELEFAX: (212) 425-7208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
US-09-794-832-9
; Sequence 9, Application US/09794832
; Patent No. 6673348
; GENERL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
; HARTL, F. Ulrich
; HOUGHTON, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION
                                                                                                                                                                               CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12 HWDFAWPW 19
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                                                                                  TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan
                                                                                                                                                                                                                                                                                                                                        - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 19;
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                                                                                                                                                                                                                                                                                                               COMPUTER NEALCH.

MEDIUM TYPER Diskette - 3. COMPUTER: 1BM compatible
COMPUTER: 1BM compatible
COMPUTER: 1BM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRIT APPLICATION DATA:
APPLICATION NUMBER: 05/002,479
FILING DATE: 13-Feb-1998
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 1995
APPLICATION NUMBER: 1995
APPLICATION NUMBER: 1996
ATTORNEY/AGENT INFORMATION:
NAME: Delaction, Richard I.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELEFAX: (212) 425-5288
INFORMATION FOR SEQ 1D NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 9; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
      HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: anino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORIGINAL «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-794-832-8; Sequence 8, Application US/09794832; Patent No. 667348; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                          ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                        NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ||||||||||
1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-011-645E-9
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Gaps
                 MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
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                                                                                                                                                                                                                                                                                - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 64; DB 2; Length 19; 100.0%; Pred. No. 0.0027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
FILING DATE: 05-0ct-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DELUCIA RICHARD L.
RECISTRATION NUMBER: 28, 839
REFERENCE/DOCKET NUMBER: 28, 839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) SEQUENCE DESCRIPTION:
) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-680-806A-8
                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-680-806A-9
; Sequence 9, Application US/09680806A
; Patent No. 6719974
; GENEBLI INFORMATION:
; APPLICANT: Sloan-Kettering Inc.
  TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 425-5288
                                                                                                                                                STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
                                                                                    NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
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Best Local Similarity 100.0
....hes 8; Conservative
                                                                                                                                                                                                                  COUNTRY: US
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        MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOB, Mee H. HOUGHTON, Alan
                                                                                                                                                                                                                                ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 64; DB 2; Length 19; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb.1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                           COMPUTER: IBM COMPATIBLE
COMPATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILLING DATE: 27-Eb-2001
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-680-806A-8
; Sequence 8, Application US/09680806A
; Patent No. 6719974
; GENERAL INFORMATION:
TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PEATURE:
OTHER INFORMATION:
                                                                                                                                                                                                                  COUNTRY: US
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Gaps
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                      MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
STREET: Cone Broadway
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                                                                                                                                                                                                                                                                   ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Length 19; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM COMPACTOR OF THE COMPUTER: IBM COMPACTOR OF COMPACTING SYSTEM: MS DOS SOFTWARE: WORD PERFECT CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/USS6/13363
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REPERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAK: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDENESS: «Unknown»
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: incernal
ORIGINAL SOURCE:
     rAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-552-868-9
; Sequence 9, Application US/09552868
; Patent No. 6761892
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: <Unknown>
                                                                                                                                                                                                  CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 HWDFAWPW 19
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TAKECHI, Yoshizumi
MAYHEW, Mark
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 10-
NG-552-868-8
NG-09-552-868-8
Sequence 8, Application US/09552868
Sequence 8, Application US/09552868
Sequence 8, Application US/09552868
SECONDARY SIOAN-Kettering Institute for Cancer Research
HARTL, FOURICH, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
                                                                                                                                                                                                                                                                 ZIF: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-Oct-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/002,479

FILING DATE: ANGUSE 18, 1995

APPLICATION NUMBER: 60/002,490

FILING DATE: ANGUSE 18, 1995

APPLICATION NUMBER: PCT/US96/13363

PILING DATE: AUGUSE 16, 1996

APPLICATION NUMBER: PCT/US96/13363

FILING DATE: PCT/US96/13363

APPLICATION NUMBER: 09/011,645

FILING DATE: PCT/US96/1363

APPLICATION NUMBER: 19, 1998

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 28, 819

REGISTRATION NUMBER: 28, 819

REGISTRATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                             CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE
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TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HARTL, F. Ulrich HOUGHTON, Alan
                                                                                                                                                                                                                                                                               - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 2; 100.0%; Pred. No. 0.0027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: PEDELUARY 13, 1998
                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PARFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILLING DATE: 10-Aug-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGBNT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REPERBUCAPOCKET NUMBER: 11746/9
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                    NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 425-5288
                                                                                                                                                                                                                               ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                 STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 HWDFAWPW 19
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US-09-636-295-9
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             MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan
                                                                                                                                                                                                                               ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 2; Length 19; 100.0%; Pred. No. 0.0027;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDENARY 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DELAUGAT, RICHARY 13, 1998
REFERENCE/DOCKET NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                        COMPUTER: IEM compatible
COMPUTER: IEM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILLING DATE: 20-Apr-2000
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHEICAL: no
FRAGNENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                   CITY: New York
STATE: NY
                                                                                                                                                                                                              COUNTRY: US
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: INDM COMPACTION TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: INDM COMPACTION COMPACTION DATA:

APPLICATION NUMBER: US/09/794,529B FILING DATE: 09-Jan-2002 CLASSIFICATION NUMBER: US/09/794,529B FILING DATE: 80-Jan-2002 CLASSIFICATION NUMBER: 00/002,479 FILING DATE: August 18, 1995 APPLICATION NUMBER: 60/002,490 FILING DATE: August 18, 1995 APPLICATION NUMBER: 09/001,496 FILING DATE: August 16, 1996 APPLICATION NUMBER: 09/011,645 FILING DATE: PEDATURY 13, 1998 ATTORNEY/AGENT INFORMATION:

NAME: DeLucia, Richard L. RESERENCE/DOCKET NUMBER: 28, 839 REFERENCE/DOCKET NUMBER: 21, 1996 ATTORNEY/AGENT INFORMATION:

TELECOMMUNICATION NUMBER: 28, 839 REFERENCE/COCKET NUMBER: 21, 1996 ATTORNEY/AGENT INFORMATION:

TELECOMMUNICATION NUMBER: 28, 839 REFERENCE/COMPUTED NUMBER: 21, 2000 ATTORNEY/AGENT NUMBER: 21, 2000 ATTORNEY/AGENT NUMBER: 21, 2000 ATTORNEY/AGENT NUMBER: 28, 839 REFERENCE/COMPUTED NUMBER: 21, 2000 ATTORNEY/AGENT NUMBER: 28, 839 ATTORNEY/AGENT NUM
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OTHER INFORMATION: hybrid peptide for human papilloma
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SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-794-529B-10
TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11. Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
RATEL, F. Ulrich
HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
                                                                                                                                                                                                                                                       CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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US-09-794-529B-11
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                                   MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
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US-09-794-529B-10
US-09-794-529B-10
Sequence 10, Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
HAPLICANT: SLOAN-Kettering Institute for Cancer Research
HARTL, F. Ulrich
HARTL, F. Ulrich
HOUGHTON, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match

100.0%; Score 64; DB 2; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.0027;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      - 3.5 inch, 1.44 Mb storage
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APPLICATION NUMBER: 60/002,479
FILING DATE: Angust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «Unknown:
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
SIREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                     NUMBER OF SEQUENCES: 30
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1 HWDFAWPW 8
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Gaps
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                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13, Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E.
                                                                                                                                                                                                                                                                                                                             ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13463
APPLICATION NUMBER: PS/US96/13463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y virus vaccine
;
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-794-529B-12
                                                                                                                                                          NUMBER OF SEQUENCES: 30
CORRESPONDENCES ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
            HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM COMPATIBLE OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 425-72 TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                           CITY: New York
                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12, Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                        ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Deflucia, Richard L.
REGISTATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION UNMERR: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                 NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: <Unknown>
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRACHENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                 CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        virus vaccine
                                                                                                                                                                                                                                                                                    COUNTRY: US
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ORGANISM: <Unknown>
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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                                                                                              TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-794-529B-14
'Sequence 14, Application US/09794529B
'Sequence 14, Application US/09794529B
'Patent No. 6641812
'GENERAL INFORMATION:
'A GENERAL INFORMATION:
'A SPPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                  3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 2;
100.0%; Pred. No. 0.0028;
tive 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIPICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA: APPLICATION NUMBER: 60/002,479
                                                                                                                                                               CORRESPONDENCE ADDRESS:
RADRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANIEM: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                         ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                             NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HWDFAWPW 8
                                                                                                                                                                                                                                                   STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-794-529B-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE
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HORTHAN, I James E.

HARTE, F. UITCH
HORE, Mean H.
HORDE, MEAN H.
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OTHER INFORMATION: hybrid peptide for human papilloma
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION NDATA:
PLING DATE: 09-Jan-2002
CLASSIFICATION NUMBER: 05/002,479
FILING DATE: APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/001,490
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 13, 1998
                                                                                                            HOUGHTON, Alan
TAKEGHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Hact Shock Protein-Based Vaccines and
Immunotherapies
                     Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus vaccine

SEQUENCE DESCRIPTION: SEQ ID NO: 16:

US-09-794-529B-16
                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
SIREET: ONE Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: <Unknown>
                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                         HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                      as
                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
ZIP: 10004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13 HWDFAWPW 20
GENERAL INFORMATION:
                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
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            ROTHMAN, James E.
HARTL, F. Ulrich
HOE, We HO.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 MD storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virus constants applied peptite sequence bescription: SEQ ID NO: 15: US-09-794-529B-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                     CITY: New York
                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
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Gaps

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RESULT 29 US-09-794-529B-17 ; Sequence 17, Application US/09794529B

RESULT 28 US-03-794-529B-16 F. Sequence 16, Application US/09794529B Patent No. 6641812

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FEATURE
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                                     APPLICANT STORM-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                    ZIF: 10004

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION UNDER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: ANGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: ANGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: ANGUST 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEBRUST 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus vaccine sequence bescription: SEQ ID NO: 17: US-09-794-529B-17
                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
Patent No. 6641812
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 HWDFAWPW 8
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                                                                                                                                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: New York
Sequence 18, Application US/09794529B
Patent No. 6641812
GENERAL INFORMATION:
ROTHMAN, James R.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: US
ZIP: 10004
COMPUTER: US
COMPUTER: DISKETCE - 3.5 inch, 1.4
MEDIUM TYPE: DISKETCE - 3.5 inch, 1.4
COMPUTER: IBM COMPATIBLE
COPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION BATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 01995
APPLICATION NUMBER: 01996
FILING DATE: AUGUST 16, 1996
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REGISTRATION NUMBER: 11746/11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFX: (Unknown)
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ylrus vaccine
seQUENCE DESCRIPTION: SEQ ID NO: 18:
US-09-794-529B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: cunknown>
TOPOLGGY: linear
MOLECTLE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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RESULT 31

RESULT 30 .US-09-794-529B-18

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TOUR TOUR TOUR TOUR TOUR TOUR TAKEN TAKEN TAKEN TAKEN TAKEN TAKEN TAKEN TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HATL, F. Ulrich HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILING DATE: 19-OCC-2001
CLASSIPICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                           Sequence 10, Application US/09794517A Patent No. 6656679 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDENDESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: New York
                        IS-09-794-517A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-794-517A-10
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                                                                                     APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TARECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529B
FILING DATE: 09-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h Similarity 100.0%; Score 64; DB 2; Length 20; Similarity 100.0%; Pred. No. 0.0028; 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: DeLucia, Richard L. REGISTRATION NUMBER: 28,839 REFERENCE/DOCKET NUMBER: 11746/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; virus vaccine
; SEQUENCE DESCRIPTION; SEQ ID NO: 19:
08-09-794-529B-19
                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
              Sequence 19, Application US/09794529B Patent No. 6641812 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 425-7200 TELEFAX: (212) 425-5288 TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                        SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                        NUMBER OF
US-09-794-529B-19
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Gaps
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                                    OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                             Query Match
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                           virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
ORGANISM: <Unknown>
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1 HWDFAWPW 8

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RESULT 34

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ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILING DATE: 19-0ct-2001
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: PCT/US96/1363
FILING DATE: PEDRIARY 13, 1998
ATTORNEY/AGENT INFERMATION: 3, 1998
ATTORNEY/AGENT INFERMATION: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                     Sequence 11, Application US/09794517A
Patent No. 665679
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
HARIL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION TELEPHONE: (212) 425-720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: NY
RESULT 33
US-09-794-517A-11
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                                                                                                                                                                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
               Sequence 12, Application US/09794517A
Patent No. 6656679
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOGHTON Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match

100.0%; Score 64; DB 2;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR AFFLIATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/ABGNT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELERX: CURLOWN:
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/794,517A FILING DATE: 19-Oct-2001 CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       yirus vaccine
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;
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
08-09-794-517A-12
                                                                                                                                                                                                                                                                                                                               Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHEICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 10004
COMPUTER READABLE FORM:
US-09-794-517A-12
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1 HWDFAWPW 8
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                                                                                                                                                                                                ROTHMAN, James E.
HARTL, F. Ulrich
HOG, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible

OPERATING SYSTEM: MS DOS

SOFTWARE: Word Perfect

CURRINT APPLICATION NUMBER: US/09/794,517A

FILING DATE: 19-Oct-2001

CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                              APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus vaccine
seQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-794-517A-13
                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                             Sequence 13, Application US/09794517A
Patent No. 6656679
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 425-5288
TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
    13 HWDFAWPW 20
                                                                  RESULT 35
US-09-794-517A-13
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                                                                                                                                                                                                             HOUGHTON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: hybrid peptide for human papilloma
RESULT 36
US-09-794-517A-14
US-09-794-517A-14
Sequence 14, Application US/09794517A
Patent No. 6656679
GENERAL INPORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, P. Ulrich
HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 677/US96/13363
FILING DATE: Adgust 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILING DATE: 19-OCt-2001
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEBRONE: (212) 425-7200
TELEFRAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
RADRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: cunknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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RESULT. 3.5
US-09-174-116
; Sequence 16, Application US/09794517A
; Patent No. 6656679
; GENERAL INFORMATION:
; ROTHWAN, James E.
HARTL, F. Ulrich
; HOUGHTON, Alan
; HOUGHTON, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1998
APPLICATION NUMBER: 09/011,645
ATORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELERX: 4UNROWN:
TELERX: 4UNROWN:
TELERX: 4UNROWN:
TELERX: 4UNROWN:
TELERX: 4UNROWN:
TELERX: 4UNROWN:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILING DATE: 19-Oct-2001
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     y virus vaccine

SEQUENCE DESCRIPTION: SEQ ID NO: 16:
US-09-794-517A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                         1 HWDFAWPW 8
                             HWDFAWPW
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                                                                                                                                                                                                                                                                                                                                                                                     ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                         Sequence 15, Application US/09794517A
Patent No. 665679
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - 3.5 inch, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4

COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WOORD PErfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILING DATE: 19-Occ-2001
CLASSIFICATION: 536
PRICA APPLICATION NUMBER: 60/002,479
FILING DATE: ANGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: ANGUST 18, 1995
APPLICATION NUMBER: PC/10956/13363
FILING DATE: ANGUST 16, 1996
APPLICATION NUMBER: PC/10956/13363
FILING DATE: ANGUST 16, 1996
APPLICATION NUMBER: PC/10956/1363
FILING DATE: PEDLUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 28,839
REPERENCE/DOCKET NUMBER: 11746/13
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SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
RADRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                            HWDFAWPW 20
HWDFAWPW 8
                                                                                                                                                                                                    US-09-794-517A-15
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Gaps ö

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Pred. No. 0.0028;
                       0; Mismatches
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STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM compatib]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (212) 425-5288
   100.04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 18
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
                     8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
Best Local Similarity
Matches 8; Conserva
                                                          1 HWDFAWPW 8
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1 HWDFAWPW
                                                                                                                                                                    -09-794-517A-18
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                                                                                                                                                                                                                         APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
FILLING DATE: 19-OCt-2001
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDFURARY 13, 1998
   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/002,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-794-517A-17
                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                    Sequence 17, Application US/09794517A Patent No. 6656679 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (212) 425-72
(212) 425-5288
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INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
 8; Conservative
                                                            13 HWDFAWPW 20
                                       1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
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   Matches
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100.0%; Score 64; DB 2; Length 20;

Query Match

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APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
IITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBRUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A
PILING DATE: 19-Oct-2001
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y virus vaccine

sequence Description: Seq ID NO: 18:
US-09-794-517A-18
                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
Sequence 18, Application US/09794517A Patent No. 665679 GENERAL INFORMATION:
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NAME: DeLucia, Richard L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                Sequence 19, Application US/09794517A
Sequence 10, Application US/09794517A
Patent No. 6656679
GENERAL INPORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
HARIL, F. Ulrich
HOGHTON, Mee H.
HOGHTON, Alan
TAKECHI, YOSHİZUMİ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517A FILING DATE: 19-Oct-2001 CLASSIFICATION: 536
100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: AAGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUSE: 1995
APPLICATION NUMBER: PCT/0996/13363
FILING DATE: AUGUSE 16, 1996
APPLICATION NUMBER: 09/011,645
APPLICATION NUMBER: 09/011,645
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/13
TELECOMMUNICATION INFORMATION:
TELEFAK: (212) 425-7200
TELEFAK: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus vaccine ;
seQUENCE DESCRIPTION: SEQ ID NO: 19: US-09-794-517A-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: One Broadway
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDENBES: «Unknown»
TOPOLLGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                  HWDFAWPW 20
                                                                                       1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 10004
                                                                                                                                                                                                                     US-09-794-517A-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James B. HARTL, F. Ulrich HOE, Mee H. HOGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM compatible
CORRETING SYSTEM: MS DOS
SOFTWARE: WOLD BY BOS
SOFTWARE: WOLD BY BOS
SOFTWARE: WOLD BY BOS
SOFTWARE: WOLD BY BOS
TILING DATE: 13-Feb-1998
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   yirus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-011-645E-10
                                                                                                                                                                                                                                                 Sequence 10, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGWENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
    Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                          1 HWDFAWPW 8
                                                                                                                        1 HWDFAWPW 8
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US-09-011-645E-10
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Gaps ö

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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                    APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                       Indels
  Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURPUTER THE DISKETTE - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WOOR Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/USS6/13363
FILING DATE: August 16, 1996
                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-011-645E-12
                                                                                                                                                                                                          Sequence 12, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
STREET: One Broadway
CITY: New York
  100.08;
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INFORMATION FOR SEQ ID NO: 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                               1 HWDFAWPW 8
                                                                                      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                         ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKEGHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                            APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIF: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 13-Feb-1998
CLASSIFICATION: 424
100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    virus vaccine;
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SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-011-645E-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                             Sequence 11, Application US/09011645E
Patent No. 6663868
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
                                                                                                            13 HWDFAWPW 20
                                                                                  1 HWDFAWPW 8
               Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FELEFAX:
                                                                                                                                                                                                            US-09-011-645E-11
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Query Match
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Length 20;

DB 2;

100.0%; Score 64;

Query Match

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virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                HWDFAWPW 8
                                                      1 HWDFAWPW 8
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                        Sequence 13, Application US/09011645E
Patent No. 6663868
GENERAL INPORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOGHTON, Alan
TAKECHI, YOSHizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION: 424
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virus vaccine sEQUENCE DESCRIPTION: SEQ ID NO: 13: US-09-011-645E-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: CUNKNOWN-
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: One Broadway CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                                                            13 HWDFAWPW 20
                                     1 HWDFAWPW 8
                                                                                                                                                           US-09-011-645E-13
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Matches
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Gaps
RESULT 46
US-09-011-645E-14
| Sequence 14, Application US/09011645E
| Patent No. 6663868
| Patent No. 6663868
| GENERAL INFORMATION:
| APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HARTL, F. Ulrich HOE, Mee H. HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi
| MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW, Mark | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | MAYHEW | M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
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APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
NAME: DELUGA, RICHARY L.
RARGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 21746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION: 424
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ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 19, 1996
ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
REFERENCE/DOCKET NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
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seQUENCE DESCRIPTION: SEQ ID NO: 16:
US-09-011-645E-16
                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                               US-09-011-645E-16
; Sequence 16, Application US/09011645E
; Patent No. 6663868
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
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STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHEICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: New York
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1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002.479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION: 424
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NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28, 839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                y virus vaccine

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SEQUENCE DESCRIPTION: SEQ ID NO: 15:

US-09-011-645E-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                          Sequence 15, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                    HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: NY
COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
                                           13 HWDFAWPW 20
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Gaps

1 HWDFAWPW 8

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JS-09-011-645E-18
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                                                                                                                                                                          APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - 3.5 inch, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 29, 8199
RESERRENCE/DOCKET NUMBER: 28,819
REGISTRATION NUMBER: 28,819
RESERRENCE/DOCKET NUMBER: 28,819
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; SEQUENCE DESCRIPTION: SEQ ID NO: 17:
08-09-011-645E-17
                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                            Sequence 17, Application US/09011645E
Patent No. 6663868
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown»
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Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
13 HWDFAWPW 20
                                                                                        JS-09-011-645E-17
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Gaps
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                                                                                                                                                                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                    APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HATL, F. Ulrich
HOE, Mee H.
HOUSHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMBUTER: IBM Compatible
CORREATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: AUGUST 18, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Deflucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKST NUMBER: 11746/1
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus vaccine

SEQUENCE DESCRIPTION: SEQ ID NO: 18:

US-09-011-645E-18
                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
Sequence 18, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULB TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
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INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
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13 HWDFAWPW 20
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JS-09-794-832-10
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                                                                                                 APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; tive 0; Mismatches 0; IndelB
                                                                                                                                                                                                                                                                                                                                                                                                               STATE: NEW TOTAL
STATE: NEW TOTAL
COUNTRY: US
ZIP: 10004
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645E
FILING DATE: 13-Feb-1998
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              yirus vaccine
;
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
08-09-011-645E-19
                                Sequence 19, Application US/09011645E Patent No. 6663868 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (212) 425-7200 (ELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon &
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWDFAWPW 8
                  US-09-011-645E-19
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                                                                                                                                                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
Sequence 10, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research ROTEMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOGHTON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM COMPATION:
COMPUTER: BM COMPATION:
SOFTWARE: WORD PERFORM
SOFTWARE: WORD PERFORM
SOFTWARE: WORD PERFORM
APPLICATION DATA:
APPLICATION NUMBER: US/09/11,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 61/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 61/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DELUCIA, RICHARD I.
REGISTRATION NUMBER: 28,839
RESPERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
RADRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Disakette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARRE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: E0/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
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                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                          Sequence 11, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28, 339
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: yea FRAGMENT TYPE: internal ORIGINAL SOURCE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                          SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                          NUMBER OF
                      JS-09-794-832-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE
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RESULT
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Gaps
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                                                                                                                                                                                                                                                TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
              Sequence 12, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOE, Mee H.
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
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100.0%; Score 64; DB 2;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 06/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
RAGISTRATION NUMBER: 28,839
REFERRENGE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECHARA: (212) 425-5288
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sequence description: SEQ ID NO: 12:
US-09-794-832-12
                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette -
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE
US-09-794-832-12
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COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MUER READABLE FORM:
COMPUTER: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARTS. WORD BETFECT CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                   HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                 Sequence 14, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDENDESS: «Unknown»
TOPOLLGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 relex: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: yes
FRAGMENT TYPE: int
ORIGINAL SOURCE:
FEATURE:
                                                                                                               US-09-794-832-14
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                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotheraples NUMBER OF SEQUENCES: 30
                                                                                                        Sequence 13, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0596/1363
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: DeLucia, Richard L.
REGISTRATION WINDER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
SIREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
    13 HWDFAWPW 20
                                                                                   US-09-794-832-13
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                       NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
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SEQUENCE DESCRIPTION: SEQ ID NO: 14:
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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                     APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James B. HARTL, P. Ulrich HOE, Mee H. HOGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - 3.5 inch, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: ET/US/05/13363
APPLICATION NUMBER: PCT/US/05/13363
APPLICATION NUMBER: ET/US/05/13363
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yirus vaccine; SEQUENCE DESCRIPTION: SEQ ID NO: 16: 16: 08-09-794-832-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                     US-09-794-832-16
Sequence 16, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                  HWDFAWPW 8
                      HWDFAWPW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE
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                                                                                                                                                                                                                             APPLICANT Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CONDTRY: US

CONDUTRY: US

ZIP: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PEFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: ET/US96/1363
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: ET/US96/1363
FILING DATE: AUGUST 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
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TELEBHOMS: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «Unknown»
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                             Sequence 15, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
                           13 HWDFAWPW 20
1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: NY
                                                                                                                   RESULT 57
US-09-794-832-15
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100.001
Best Local Similarity 100.
Matches 8; Conservative
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ZIP: 10004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NY
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1 HWDFAWPW
                                                          1 HWDFAWPW
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM comparible
COMPUTER: WAR DOS
SOFTWARE: WAR PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                         APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
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SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: DeLucia, Richard L.
                                                                                                                                                                    Sequence 17, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 425-72
TELEFAX: (212) 425-5288
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INFORMATION FOR SEQ 1D NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCES:
                                                                          13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
                                       1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF
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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOE, HOE H. HOUGHTON, Alan TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                - 3.5 inch, 1.44 Mb storage
  Indels
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FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILLING DATE: 27-Peb-2001
CLASSIFICATION: «UNKNOWN»
PRIOR APPLICATION DATA:
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEPAX: (212) 425-528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus vaccine

SEQUENCE DESCRIPTION: SEQ ID NO: 18:
US-09-794-832-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                 Sequence 18, Application US/09794832
Patent No. 6673348
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: <URNOWN>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION
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INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                         MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM
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                                                                                                                                                                                                                                                                                               APPLICANT: SLOAD-Kettering Institute for Cancer Research ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown>
PRIOR APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: E0/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
TELEREBERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; virus vaccine; sEQUENCE DESCRIPTION: SEQ ID NO: 19: US-09-794-832-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                         Sequence 19, Application US/09794832; Patent No. 6673348; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 19
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
  Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                   HWDFAWPW 20
                                                                                        1 HWDFAWPW 8
                                                                                                                                                                                                                    US-09-794-832-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE
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US-09-680-806A-10
19-09-680-806A-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
COMPUTER: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM COMPALLAL
COMPUTER: WAS DOS
SOFTWARE: WATCH PERFOCE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-Oct-2000
CLASSIPICATION A24
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 61/995
FILING DATE: AUGUST 18, 1995
FILING DATE: AUGUST 16, 1996
FILING DATE: AUGUST 16, 1996
FILING DATE: AUGUST 16, 1996
FILING DATE: AUGUST 16, 1996
FILING DATE: AUGUST 16, 1996
FILING DATE: AUGUST 16, 1998
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NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHEICAL: yes
FRAGWENT TYPE: internal
ORIGINAL SOURCE:
    Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                         1 HWDFAWPW 8
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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James B.
HARTL, P. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                  100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0996/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/680,806A FILING DATE: 05-Oct-2000 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
TELEFHONE: (212) 425-7200
                                                                                                             0; Mismatches
; SEQUENCE DESCRIPTION: SEQ ID NO: 11: US-09-680-806A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                             US-09-680-806A-12; Sequence 12, Application US/09680806A; Batent No. 6719974; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PETECT
CURRENT APPLICATION DATA:
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STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 12
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 30
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                 Query Match
Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible COMPUTER: IBM compatible SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/680,806A FILLING DATE: 05-0ct-2000 CLASSIFICATION: 424 PRIOR APPLICATION DATA:
                                                 Length 20;
                                                 100.0%; Score 64; DB 2; 100.0%; Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
PRILING DATE: August 18, 1995
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13163
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                        Sequence 11. Application US/09680806A Patent No. 6719974 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 11.
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
                                                                                        Conservative
                              13 HWDFAWPW 20
                                                                                                                                 1 HWDFAWPW
        US-09-680-806A-10
                                                                                                                                                                                                                                                         US-09-680-806A-11
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us-10-053-520-143.rai

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OTHER INFORMATION: hybrid peptide for human papilloma
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HARTL, F. Ulrich
HOG, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - 3.5 inch, 1.44 Mb storage
                                                                                                                                                             0; Indels
                                                                                                            Query Match 100.0%; Score 64; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4.

COMPUTER: IBM compatible
CORREATING SYSTEM: MS DOS
SOFTWARE: WOLD PETFOCT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: OS-Oct-200
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDTUATY 13, 1998
ATTORNEY AGENT INFORMATION:
NAWE: DELUCAT ON NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
                        virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                            RESULT 66
US-09-680-806A-14
Sequence 14, Application US/09680806A
Statent No. 6719974
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: One Broadway CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGWENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
                                                                                                                                                                                                                                        1 HWDFAWPW 8
                                                                 US-09-680-806A-13
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US-09-806-806A-13
Sequence 13, Application US/0968086A
Sequence 13, Application US/09680806A
Sequence 13, Application US/09680806A
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHYOW, Alan
HOUGHYOW, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
MAYHEW, Mark
HARE Shock Protein-Based Vaccines and
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATION SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRINT APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-0cc-2000
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/001,490
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: AUGUST 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies NUMBER OF SEQUENCES: 30
                                                                                           Length 20;
                                                                                         Query Match 100.0%; Score 64; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
    ; virus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-680-806A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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                                        OTHER INFORMATION: hybrid peptide for human papilloma
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                        Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
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APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-0ct-2000
CLASSIFICATION 1424

PRIOR APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0896/13363
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: Ebbruary 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 15:
US-09-680-806A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                               virus vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                                                                                                                                                                                                           1 HWDFAWPW 8
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                   OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOG; Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                     Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: OS-0Ct-2000
CLASSIFICATION: 424
PRIOR APPLICATION: A24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                    virus vaccine
;
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-09-680-806A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                    US-09-680-806A-15

US-09-680-806A-15

Sequence 15, Application US/09680806A

Patent No. 6719974

GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
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ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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STREET: One Broadway
CITY: New York
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FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide
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1 HWDFAWPW 8
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                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
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US-09-680-806A-18
                                                            FEATURE:
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                                                              OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARIL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TARECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                 Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-0ct-2000
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION:
                                                                                    virus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO: 16:
US-09-680-806A-16
                                                                                                                                                                                                                                                                                                                                                                            US-09-680-806A-17
; Sequence 17, Application US/09680806A
; Patent No. 6719974
; GENERAL INFORMATION:
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STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 17
SEQUENCE CHARACTERISTICS
  ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                             13 HWDFAWPW 20
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OTHER INFORMATION: hybrid peptide for human papilloma
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ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
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Similarity 100.0%; Score 64; DB 2; Length 20;
Similarity 100.0%; Pred. No. 0.0028;
8; Conservative 0; Mismatches 0; Indels
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APPLICATION NUMBER: 60/002,479
FILING DATE: ANGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: ANGUSE 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: ANGUSE 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBRUSER: 09/011,645
FILING DATE: FEBRUSER: 13, 1998
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OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-Oct-2000
CLASSIFICATION: 424
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NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
TELECOMMUNICATION INFORMATION:
                     y virus vaccine; SEQUENCE DESCRIPTION: SEQ ID NO: 17: US-09-680-806A-17
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ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18, Application US/09680806A Patent No. 6719974 GENERAL INFORMATION:
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TELEX: <Unknown>
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STRANDEDNESS: <Unknown>
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Gaps
                                                                                           ) OTHER INFORMATION: hybrid peptide for human papilloma; sequence describe; SEQUENCE DESCRIPTION; SEQ ID NO: 19:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James E.
HARTL, F. Ulrich
HOBE, Mee H.
HOUGHIYON, Alan
TARECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                  100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 90/011,645
FILING DATE: FEBRUARY 13, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM; MS DOS
SOPTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
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STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
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LENGTH: 20
                       HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.4
....neg 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                  1 HWDFAWPW 8
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US-09-800-806A-19
i Sequence 19, Application US/09680806A
j Patent No. 6719974
i GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
KOTHANN, James E.
HARTL, F. Ulrich
HOUG: Mee H.
HOUG: Mee H.
HOUGHOW, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
MAYHEW, Mark
HOUGHOW, Mark
                                                                                                                                                                                                                                                                            ;
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                                                                                                              OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                              Score 64; DB 2; Length 20;
Pred. No. 0.0028;
                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1955
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0896/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBLUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,806A
FILING DATE: 05-Oct-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/10
                                                                                                                                      ; virus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO: 18:
US-09-680-806A-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.0%;
Matches 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS: LENGTH: 20
                     FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: New York
HYPOTHETICAL: yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
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                                                                                        FEATURE:
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OTHER INFORMATION: hybrid peptide for human papilloma
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER IBM compatible COMPUTER: IBM compatible COMPUTER: IBM compatible COMPUTER: IBM compatible COMPUTER: IBM compatible COMPUTER: Ward Perfect CURRENT APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: August 16, 1996
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: August 16, 1996
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 20,0101,645
FILING DATE: August 18, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REGISTRATION NUMBER: 28,839
REGISTRATION NUMBER: 28,839
                                                                                                                                                                                                                                                                               Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                   virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION
                      TOPOLOGY: linear
MOLECULB TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <UNKNOWT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: New York
STATE: NY
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                                                                                                                                                      OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HATL, F. V. Urich
HOE, Mee H.
HOKHTHON, Alan
TAKEHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                             0: Indels
                                                                                                                                                                                                                                                                Query Match 100.0%; Score 64; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBRUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
                                                                                                                                                                                 virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-552-868-11
; Sequence 11, Application US/09552868
; Patent No. 6761892
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
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TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: YES
FRACKENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
STATE: NY
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                                                                                                                                  FEATURE:
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Gaps

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OTHER INFORMATION: hybrid peptide for human papilloma
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ROTHWAN, James E.

HARTL, F. Ulrich
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
IITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                      100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                0; Indels
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COMPUTER: IBM compatible
OPERATING SYSTEM: NS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/52,868
FILING DATE: 20-Apr-2000
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
APPLICATION NUMBER: PCT/US96/13163
                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: DeLucia, Richard L. REGISTRATION WUMBER: 28,639
REFERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                 virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
TYPE: amino acid
STRANDENESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 14, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                             ORGANISM: <Unknown:
                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 8; Conservative
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US-09-552-868-14
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                                                                                                                                                                                  OTHER INFORMATION: hybrid peptide for human papilloma
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ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - 3.5 inch, 1.44 Mb storage
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FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
FILING DATE: FEbruary 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                          virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 13, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Kenyon & Ke
STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                        1 HWDFAWPW 8
                                                                                                                                                             FEATURE:
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Gaps

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APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MATHEN, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
RADDRESSEE Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
ODERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-200
CLASSIFICATION: 424
PRIOR APPLICATION HATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/8
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                        virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
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   INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
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ZIP: 10004
COMPUTER READABLE FORM:
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 HWDFAWPW 8
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US-09-552-868-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: STATE APPLICANT: STATE APPLICANT: STATE APPLICANT: STATE APPLICANT: And BE HARTL, F. Ulrich HOE, Mee HOUGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and MANDEN OF CHARLES APPLICANT AND APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
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APPLICATION NUMBER: 60/002,479
FILING DATE: Angust 18, 1955
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/0956/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBRUARY 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM compatible
OPBRAITING SYSTEM; MS DOS
SOFTWARB: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
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SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
SEQUENCE CHARACTERISTICS:
LENGTH: 20
TYPE: amino acid
STRANDEDNESS: cUnknown>
TOPOLOGY: linear
WOLECTLE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-552-868-15; Sequence 15, Application US/09552868; Sequence 15, Application US/09552868; Patent No. 6761892; GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: One Broadway
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                                                                                                                                                                                                                                                                                                                                                 PEATURE
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich HOE, Mee H. HOUE, Mee H. HOUETHON, Alan TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: IBM compatible
SOFTHARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: PET/0896/13363
FILING DATE: AUGUSE 16, 1996
APPLICATION NUMBER: PET/1996
PILING DATE: EPDLUALY 13, 1998
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REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                      tirus vaccine sequence DESCRIPTION: SEQ ID NO: 17: US-09-552-868-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                            TYPE: amino acid
STRANDENESS: «Unknown»
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION
                     INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: New York
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US-09-552-868-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TARECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/52,868
FILING DATE: 20-Apr-2000
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1995
APPLICATION NUMBER: PCT/US96/1363
FILING DATE: August 16, 1996
APPLICATION NUMBER: PCT/US96/1363
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REFERENCE/DOCKET NUMBER: 11746/8
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                          virus vaccine
sEQUENCE DESCRIPTION: SEQ ID NO: 16:
US-09-552-868-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                  TYPE: amino acid
STRANDENESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
              INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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CITY: New York
                                                                                                                                                                                                                                             ORGANISM: <Unknown>
TELEX: <Unknown>
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1 HWDFAWPW 8
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
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COMPUTER: 1BM compatible
COMPUTER: 1BM compatible
SSTATING SYSTEM: MS DOS
SOFTMARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/552,868
FILING DATE: 20-Apr-2000
CLASSIFICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 18, 1996
APPLICATION NUMBER: 09/011,645
PILING DATE: ROBUST 13, 1998
ATTORNEY AGENT INFORMATION:
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REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine seculos seq ID NO: 18: US-09-552-868-18
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19, Application US/09552868
Patent No. 6761892
GENERAL INFORMATION:
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US-09-552-868-19
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TELEPRONICATION INTROMATION:
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TELEPRONICATION OF 15:
STRANDENESS: "CUMCOON-)
TELEPRONICATION: ON 15:
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Gaps
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HOE, Mee H.
HOUGHTON, Alan
TAKEGHI, YOSHİZUMİ
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E.
HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               - 3.5 inch, 1.44 Mb storage
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APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 90/011,645
ATTORNEY/AGENT INFORMATION: 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM compatible
OPERATING SYSTEM; MS DOS
OPERATING SYSTEM
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10 - Aug-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                               virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                              TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                            ORGANISM: <Unknown>
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ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
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ZIP: 10004

COMPUTER READABLE FORM:
COMPUTER: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word PATECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
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FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
REFERENCE/DOCKET NUMBER: 11746/9
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SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                  TYPE: amino acid
STRANDENESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
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NAME: DeLucia, Richard L
                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 HWDFAWPW 20
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US-09-636-295-11
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TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - 3.5 inch, 1.44 Mb storage
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APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-ANG-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                        virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
                                                                                                                                                                                                                LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 16, 1996
APPLICATION NUMBER: 60/001,490
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13, Application US/09636295

Patent No. 6773707

GENERAL INPORMATION:
APPLICANT: SIOAn-Kettering Institute for Cancer Research ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028;
             REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWIN»
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                           TYPE: amino acid
STRANDBDNESS: «Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 HWDFAWPW 8
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OTHER INFORMATION: hybrid peptide for human papilloma
NAME: DELUCIA, RICHARD L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: (UNKNOWN)
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
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Gape

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NY
COUNTRY: US
ZIP: 10004
COMPUTER: READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compactible
COMPUTER: IBM compactible
CORRATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HATTL, F. Ulrich
HOE, Mee H.
HOKE, Mee H.
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 2; Length 20; 100.0%; Pred. No. 0.0028; rive 0; Mismatches 0; Indels
                     FILING DATE: February 13, 1998
ATTORNEY AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMONICATION INFORMATION:
TELEPHONE: (212) 425-5208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE DESCRIPTION: SEQ ID NO: 15: US-09-636-295-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                  TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: August
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               virus vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HWDFAWPW 8
                                                                                                                                                                                                                                                                           LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIPICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: COT/US96/13363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT/US96/13363
16, 1996
                  ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 14:
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15, Application US/09636295
Patent No. 6773707
GENERAL INFORMATION:
                                                                                                                                                   TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: August
                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13 HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE
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APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
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sEQUENCE DESCRIPTION: SEQ ID NO: 17:
US-09-636-295-17
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Patent No. 6773707
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLLGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & J
STREET: One Broadway
                                                                                                                                                                                                                              TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                               LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 HWDFAWPW 8
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COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compactible
COMPUTER: IBM compactible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 64; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 0.0028; Matches 8; Conservative 0; Mismatches 0; Indels
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard 1.
REGISTRATION NUMBER: 28,839
REGISTRATION NUMBER: 11746/9
TELEPRONICATION INFORMATION:
TELEPHONE: (212) 425-5288
TELEFAX: (212) 425-5288
TELEX: <UNKNOWN>
INFORMATION POR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STRRET: One Broadway
                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 17, Application US/09636295; Patent No. 6773707; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 HWDPAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 64; DB 2; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: FEBRUARY 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DELUCIA, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
```

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich HOB, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark

- 3.5 inch, 1.44 Mb storage

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490

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Search completed: March 24, 2006, 12:35:57 Job. time : 46 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
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; Sequence 19, Application US/09636295
; Sequence 19, Application US/09636295
; GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
HARTIL F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshiumi
MAYHEW, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies

NUMBER OF SEQUENCES: 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: hybrid peptide for human papilloma
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: Ibm compatible COMPUTER: Ibm compatible COMPUTER: SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATE: US/109/636,295
FILING DATE: 10-Aug-2000
CLASSIFICATION: 424
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FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/ABGNT INFORMATION:
NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELERX: 4UNLOWN:
TELERX: 4UNLOWN:
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
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SEQUENCE DESCRIPTION: SEQ ID NO: 18:
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ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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ZIP: 10004
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Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 8; Conservative 0; Mismatches 0; Indels
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: DELUCIA, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/9
TELECOMMUNICATION:
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NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
ZOUTRY: US
ZONPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                              - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: heat shock binding sequence SEQUENCE DESCRIPTION: SEQ ID NO: 1:
COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 "...

COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: 09/011,645
FILING DATE: AUGUSE: 09/011,645
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FILING DATE: AUGUSE: 09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,09/01,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: mouse
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Publication No. US20030021794A1
GENERAL INFORMATION:
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mae
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunotherapies
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CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
CONTEX: NY
ZIP: 10004
                US-09-794-517-1
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Gaps

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RESULT 2
US-09-794-529-1
Sequence 1, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
HAPLICANT: Sloan-Kettering Institute for Cancer Research
HARTL, F. Ulrich
HARTL, F. Ulrich
HOG, Mee H.
HOGHTON, Alan
THOUGHTON, Alan
THOUGHTON, Alan
THOUGHTON, Alan
THOUGHTON, Alan
THOUGHTON, Heat Shock Protein-Based Vaccines and
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Gaps
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Publication No. US20030134787A1

GENERAL INFORMATION:

APPLICANT: Sloan-Kettering Institute for Cancer Research

APPLICANT: Rothman, James E.

APPLICANT: Mayhew, Mark

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Morefelli, Ouathek

APPLICANT: Moroi, Yolchi

TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES

FILE REFERENCE: 11746/46003

CURRENT APPLICATION NUMBER: US/10/052,578

FRIOR APPLICATION NUMBER: 08/961,707

PRIOR APPLICATION NUMBER: 08/961,707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 64; DB 3; Length 8; 100.0%; Pred. No. 1.7e+06;
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                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-F60-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: AUGUST 16, 1996
ATTORNEY/AGRNT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                             APPLICATION NUMBER: US/09/794,832 FILING DATE: 27-Feb-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 321
SOFTWARE: WordPerfect 8.0 for Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION TELEPHONE: (212) 425-720
                         OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: mouse
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Best Local Similarity 100.
Matches 8; Conservative
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Publication No. US20030082198A1
GENERAL INFORMATION:
APPLICANT: SOTHMAN, James E.
HARTL, F. Ulrich
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 64; DB 3; Length 8; Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     heat shock binding sequence
                                                                                                                                                                                                              FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
                                                                                          APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                    PRIOR APPLICATION DATA: APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: heat shock
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 425-5288
TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECTLE TYPE: peptide
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ADDRESSEE: Kenyon & STREET: One Broadway
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYPOTHETICAL: no FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: mouse
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COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HWDFAWPW 8
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TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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                   COMPUTER READABLE FORM:
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STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HWDFAWPW 8
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US-10-053-520-143

US-10-053-520-143

Sequence 143, Application US/10053520

Publication No. US2030166530A1

GENERAL INFORMATION:

APPLICANT: Sloan-Kettering Institute for Cancer Research

APPLICANT: Rochman, James E.

APPLICANT: Hoe, Mee H.

APPLICANT: Houghcon, Alan

APPLICANT: Houghcon, Alan

APPLICANT: Houghcon, Alan

APPLICANT: Worfell!, Outhek

TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES

FILE REPERENCE: 11746/46004

CURRENT APPLICATION NUMBER: US/10/053,520

CURRENT FILING DATE: 2002-10-01

PRIOR FILING DATE: 1997-10-31

NUMBER OF SEQ ID NOS: 321

SOFTWARE: WordPerfect 8.0 for Windows

SEQ ID NO 143

LENGTH: 8

TUTLE OF INVENTION NUMBER: OF OF WINDOWS

SEQ ID NO 143
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TAKECHI, Yoshizumi
MAYHEW, Mark
IITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOE, Mee H.
                                                                                                    Query Match 100.0%; Score 64; DB 4; Length 8; Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; I
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
JOTHER INFORMATION: peptide in ml3 coliphage
US-10-053-520-143
                                     ; OTHER INFORMATION: peptide in ml3 coliphage US-10-052-578-143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunotherapies
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ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-170-713A-1
; Sequence 1, Application US/10170713A
; Publication No. US20030185842A1
; GENERAL INFORMATION:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 30
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                                                                                                                                                                                                                    1 HWDFAWPW 8
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US-10-71-734-1
; Sequence 1, Application US/10171734
; Publication No. US20030185843A1
; GENERAL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
; TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
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CUMPUTER: EDISKette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: 13-Jun-2002
CLASSIFICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 09/011,645
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 16, 1996
ATTORNEY/AGENT INPORMATION:
NAME: CASSON, Lawrence P
REGISTRATION NUMBER: 46,606
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Best Local Similarity 100.0%; Score 64; DB 4; Length 8;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: heat shock binding sequence SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
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                                                                                                                                                                                                                                                                                                 0; Indels
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Publication No. US20040052812A1

GENERAL INFORMATION:

APPLICANT: Mee Hoe

TITLE OF INVENTION: VACCINES

TITLE OF INVENTION: VACCINES

TITLE OF INVENTION WHERE: US/10/258,146A

CURRENT APPLICATION NUMBER: US/10/258,146A

CURRENT APPLICATION NUMBER: 0S/10/258,146A

PRIOR APPLICATION NUMBER: 0S/10/256B

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 1800

SOSTWARE: FRASESQ for Windows Version 3.0
                                                                                                                                                                                                                                                Query Match
100.0%; Score 64; DB 4; I
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rothman, James
APPLICANT: Rothman, James
APPLICANT: Hoe, Met
TILE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT APPLICATION NUMBER: 60/197,462
PRIOR APPLICATION NUMBER: 60/197,462
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 8
                                                                                                                                                ; FEATURE:
; OTHER INFORMATION: peptide in m13 coliphage
US-10-053-498B-143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: insert in M13 coliphage US-10-258-147-3
NUMBER OF SEQ ID NOS: 321
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 3, Application US/10258147
Publication No. US20040043419A1
GENERAL INPORMATION:
                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Aritifical Sequence
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1 HWDFAWPW 8
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1 HWDFAWPW 8
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APPLICANT: Rothman, James E.
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Hartl, Ulrich
APPLICANT: Morol, Yolchi
APPLICANT: Morol, Yolchi
TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
FILE REPERENCE: 11746/460002
CURRENT APPLICATION NUMBER: 2002-01-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PEATURE:
OTHER INFORMATION: heat shock binding sequence SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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100.0%; Score 64; DB 4; I
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDRUARY 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Casson, Lawrence P.
REGISTRATION UNDHER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                   APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIPICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 143, Application US/10053498B Publication No. US20030194409A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2002-01-17
PRIOR APPLICATION NUMBER: 08/961,707
PRIOR FILING DATE: 1997-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 HWDFAWPW 8
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1 HWDFAWPW 8
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US-10-367-594-1
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                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/10328953

| Sequence 3, Application US/10328953
| Publication No. US20040071656A1
| GENERAL INFORMATION:
| APPLICANT: Wieland, Felix
| APPLICANT: Wieland, Pelix
| TITLE OF INVENTYON: Modulation of Heat-Shock-Protein-Based Immunotherapies
| TITLE OF INVENTYON: Modulation of Heat-Shock-Protein-Based Immunotherapies
| FILE REFRENCE: 11390/46.01
| CURRENT FILING DATE: 2002-12-23
| PRIOR APPLICATION NUMBER: US 60/342,570
| PRIOR APPLICATION NUMBER: US 60/342,570
| PRIOR PILING DATE: 2001-12-28
| PRIOR PILING DATE: 2001-12-28
| PRIOR PILING DATE: 2002-04-12
| PRIOR PILING DATE: 2002-09-28
| PRIOR PILING DATE: 2002-09-28
| PRIOR PILING DATE: 2002-09-28
| PRIOR PILING DATE: 2002-09-28
| NUMBER OF SEQ ID NOS: 331
| SOFTWARE: WordPerfect 8.0 for Windows SEQ ID NO 3
| LENGTH 8
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APPLICANT: Rothman, James E.

APPLICANT: Harl, F. Ulrich

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

TITLE OF INVENTION WIMBER: US, 10/367, 580

CURRENT APPLICATION NUMBER: US, 09/794, 832

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1996-08-16
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                                                                                     Length 8;
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                                                                               Query Match
100.0%; Score 64; DB 4; L
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0;
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100.0%; Score 64; DB 4; I
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: synthetic peptide US-10-328-953-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/10367580 Publication No. US20040071720A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                            1 HWDFAWPW 8
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1 HWDFAWPW 8
; ORGANISM: human
US-10-258-146A-1
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APPLICANT: Rochman, James E.
APPLICANT: Rochman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11146/46.012
CURRENT APPLICATION NUMBER: US/10/367,593
CURRENT APPLICATION NUMBER: US 09/011,645
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
NUMBER OF SEQ ID NOS: 349
NUMBER WordPerfect 8.0 for Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 1
LENGTH: 8
                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: synthetic peptide US-10-367-580-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: synthetic peptide US-10-367-593-1
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                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDPAWPW 8
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION WHBER: US/10/367,654
CURRENT FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 99/636,295
PRIOR PILING DATE: 2000-08-10
PRIOR PILING DATE: 1996-08-16
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
THE REFERRENCE: 17746/461041
CURRENT APPLICATION NUMBER: 02(1/0)367,594
CURRENT FILING DATE: 2003-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 64; DB 4; Length 8; Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                        CURRENT FILING DATE: 2003-02-14.
PRIOR FILING DATE: 2000-10-05
PRIOR PILING DATE: 2000-10-05
PRIOR PILING DATE: 1900-10-05
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows SEQ ID NO 1
ERROR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows SEQ ID NO 1
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SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 1
LENGTH: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10367654 Publication No. US20040071723A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Seguence
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APPLICANT: Prince-Cohane, Kenya
APPLICANT: Prince-Cohane, Kenya
APPLICANT: Brince-Cohane, Kenya
APPLICANT: Mehta, Sunil
APPLICANT: Andjelic, Sofija
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Barber, Brian
APPLICANT: Andjelic, Sofija
APPLICANT: ANGROUED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF INVENTION: IMPROVED: 2004-09-99
CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 8; 100.0%; Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
PRIOR APPLICATION NUMBER: US 10/170,738
PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 09/552,868
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 1996-02-13
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
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PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WORDERFECT 8.0 for Windows
SEQ ID NO 1
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 358
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APPLICANT: Fletchner, Jessica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTHER INFORMATION: synthetic peptide US-10-367-674-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence FEATURE:
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Best Local Similarity 100.C
Matches 8; Conservative
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Matches 8; Conserv
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hose Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: Hower, 1746/4610211
CURRENT APPLICATION NUMBER: US(110/367,674
CURRENT FILING DATE: 2003-02-14
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Oais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTORIN MEDIATED INVUNCTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Score 64; DB 4; Length 8;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels
        PRIOR FILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: synthetic peptide US-10-367-668-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 36, Application US/10258144; Publication No. US20040101532A1
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ORGANISM: Artificial Sequence
FEATURE:
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1 HWDFAWPW 8
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Length 8;
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Pred. No. 1.7e+06;
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APPLICANT: Frank Landsberger

TITLE OF INVENTION: HEAT SHOCK PROTEIN-BASED ANTIVIRAL

TITLE OF INVENTION: VACCINES

FILE REFERENCE: 11390/46301

CURRENT APPLICATION NUMBER: US/10/258,146A

CURRENT FILING DATE: 2003-09-05

PRIOR APPLICATION NUMBER: PCT/US01/12568

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR PILING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 180

SOFTWARE: PASELSEQ for Windows Version 3.0

SEQ ID NO 175
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TITLE OF INVENTION: HEAT SHOCK PROTEIN-BASED ANTIVIRAL
TITLE OF INVENTION: VACCINES
FILE REFERENCE: 11390/46301
CURRENT APPLICATION NUMBER: US/10/258,146A
CURRENT FILING DATE: 2003-09-05
FRIOR APPLICATION NUMBER: PCT/US01/12568
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR APPLICATION NUMBER: 0190,462

                                    1.7e+06;
100.0%; Score 64; DB 5; 100.0%; Pred. No. 1.7e+06
                                                                        0; Mismatches
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; Sequence 176, Application US/10258146A
; Publication No. US20040052812A1
; GENERAL INFORMATION:
; APPLICANT: Mee Hoe
                                                                                                                                                                                                                                                                                                                                                                            Sequence 175, Application US/10258146A Publication No. US20040052812A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
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       Query Match
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| GENERAL INFORMATION: | APPLICANT: Pictuce-Cohane, K. |
| APPLICANT: Prince-Cohane, K. |
| APPLICANT: Prince-Cohane, K. |
| APPLICANT: Mehta, S. |
| APPLICANT: Mehta, S. |
| APPLICANT: Andjelic, S. |
| APPLICANT: Andjelic, S. |
| APPLICANT: Andjelic, S. |
| APPLICANT: Andjelic, S. |
| APPLICANT: Barber, BROVED HEAT SHOCK PROTEIN-BASED TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED TITLE OF INVENTION: UNGER: US/10/820,067A |
| FILE REFERENCE: 8449-406-999 |
| CURRENT APPLICATION NUMBER: US/40/82,469 |
| PRIOR FILING DATE: 2003-04-11 |
| PRIOR PELICATION NUMBER: 60/463,746 |
| PRIOR APPLICATION NUMBER: 60/503,417 |
| PRIOR APPLICATION NUMBER: 60/503,417 |
| PRIOR FILING DATE: 2003-09-16 |
| SOFTWARE: FastSEQ for Windowe Version 4.0 |
| SECTION 10 069
                                                                                                                                                                                                                                                            APPLICANT: Menca. S.
APPLICANT: Menca. S.
APPLICANT: Andjelic, S.
APPLICANT: Andjelic, S.
TITLE CANT: Barber, B.
TITLE COP INVENTION: INPROVED HEAT SHOCK PROTEIN-BASED
TITLE OF INVENTION: VACCINES AND IMMUNOTHERAPIES
FILE REPERBENCE: 8449-466-999
CURRENT APPLICATION NUMBER: 05/10/820,067A
CURRENT FILING DATE: 2004-04-08
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR PAPLICATION NUMBER: 60/463,746
PRIOR PAPLICATION NUMBER: 60/503,417
PRIOR PLING DATE: 2003-04-18
PRIOR PLING DATE: 2003-09-16
NUMBER OF SEQ ID NOS: 926
SOFTWARE: PASCESEQ for Windows Version 4.0
LENGTH: B
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US-10-820-067A-267
                                                                    Application US/10820067A
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ORGANISM: Artificial Sequence
                                                                                                        US20050214312A1
                                                                                                                                                                      APPLICANT: Fletchner, J.
APPLICANT: Prince-Cohane, K.
APPLICANT: Mehta, S.
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US-10-820-067A-869
                                                                    Sequence 267, Ag
Publication No.
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APPLICANT:
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46101
CURRENT FILING DATE: 2002-10-17
PRIOR PRICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FRAESEQ for Windows Version 3.0
SEQ ID NO 278
LENGTH: 16
                                                                                                                                                                                                                                                   APPLICANT: Livinget, This APPLICANT: Livinget, This APPLICANT: Al-Awgati, Qais APPLICANT: Al-Awgati, Qais APPLICANT: Al-Awgati, Qais APPLICANT: How, Mack APPLICANT: Hoe, Mee TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA FILE REFERENCE: 11746/46401 CURRENT APPLICATION NUMBER: US/10/258,144 FRICK APPLICATION NUMBER: 60/197,642 PRICK APPLICATION DATE: 2000-04-17
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100.0%; Score 64; DB 4; Length 16;
Best Local Similarity 100.0%; Pred. No. 0.059;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 277
LENGTH: 16
                                                                                                RESULT 27
US-10-258-144-277
Sequence 277, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
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Publication No. US20040101532A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens derivative
US-10-258-144-277
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ORGANISM: Homo sapiens derivative
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1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: obtained from a phage synthetic peptide library US-10-258-146A-172
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                                                                                                                                                                US-10-258-146A-171

Sequence 171, Application US/10258146A

Publication No. US20040052812A1

Publication No. US20040052812A1

GENERAL INFORMATION:

APPLICANT: Frank Landsberger

TITLE OF INVENTION: HEAT SHOCK PROTEIN-BASED ANTIVIRAL

TITLE OF INVENTION: WACCINES

FILE REPERENCE: 11390/46301

CURRENT APPLICATION NUMBER: US/10/258,146A

CURRENT FILING DATE: 2003-09-05

PRIOR APPLICATION NUMBER: ECT/US01/12568

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 180

SOFTWARE: PastSEQ for Windows Version 3.0

SEQ ID NO 171

LENGTH: 12
  0; Indels
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APPLICANT: Mee Hoe

APPLICANT: Frank Landsberger

TITLE OF INVENTION: HEAT SHOCK PROTEIN-BASED ANTIVIRAL

TITLE OF INVENTION: WACCINES

FILE REFERENCE: 11390/46301

CURRENT FILING DATE: 2003-09-05

PRIOR APPLICATION NUMBER: PCT/US01/12568

PRIOR APPLICATION NUMBER: PCT/US01/12568

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR FILING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 180

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 172

LENGTH: 12
  0; Mismatches
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  8; Conservative
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagari, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTMARE: PRESESCE for Windows Version 3.0
SSOFTMARE: PRESESCE for Windows Version 3.0
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                                                                   APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
EURRENT APPLICATION NUMBER: 60/202-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hos, Mee
APPLICANT: Hos, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
ENG ID NO 217
LENGTH: 17
TYPE: PRT
ORGANISM: Homo sapiens derivative
US-10-258-1144-217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 218, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
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Livingston, Phil
                       Al-Awgati, Qais
Mayhew, Mark
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APPLICANT: Houghton, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Me
TTITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: MABER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR PELICATION NUMBER: 60/197,642
PRIOR PELICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 208
LENGTH: 17
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               Sequence 208, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
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Publication No. US20040101532A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens derivative US-10-258-144-209
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Matches 8; Conservative
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Best Local Similarity
Matches 8; Conserv
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                                                                                                               TYPE: PRT
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERENCE: 11746/446101
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 0002-10-17
PRIOR PRILING DATE: 2000-04-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 228
LENGTH: 17
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US-10-259.
US-10-259.

Sequence 237, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17
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TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA PILE REPERBURE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PELLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FRRUESQ for Windows Version 3.0
LENGTH: 17
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.062;
rative 0; Mismatches 0; Indels
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PRIOR APPLICATION NUMBER: 60/197,642
                                                                                                                                                                                                                     ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-227
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Best Local Similarity 100.1
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT PELICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-04-17
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Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
  100.0%; Pred. No. 0.062;
tive 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 258
LENGTH: 17
TYPE: PRT
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 287
LENGTH: 17
TYPE: PRT
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                Sequence 258, Application US/10258144; Publication No. US20040101532A1
GENERAL INFORMATION:
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Best Local Similarity 100. Matches 8; Conservative
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1 HWDFAWPW 8
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awqazi, Qais
APPLICANT: Al-Awqazi, Qais
APPLICANT: Al-Awqazi, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
FRICR APPLICATION NUMBER: 60/197,642
FRICR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 257
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                          Sequence 248, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Hayingston, Phil

APPLICANT: Mayhew, Mark

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTHARE: FastESQ for Windows Version 3.0
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                                                                                                 Query Match
100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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; TYPE: PRT ; ORGANISM: Homo sapiens derivative US-10-258-144-247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-248
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LENGTH: 17
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Sequence 298, Application US/10258144

Publication No. US20040101532A1

GENERAL INPORMATION:
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR PLICATION NUMBER: 60/197,642
PRIOR PLICATION NUMBER: 60/197,642
PRIOR PLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOCTHARE: PSECSOF OF Windows Version 3.0
SEQ ID NO 298
LENGTH: 17
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Publication No. US20040101532A1

GENERAL INFORMATION.

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awagati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Hoe, Me

TITLE OF INVENTION METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2000-210-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR PLING DATE: 2000-04-17

WUMBER OF SEQ ID NOS: 502

SOFTMARE: FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEE FREEEEE FREEEE                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens derivative US-10-258-144-298
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Best Local Similarity 100.
Matches 8; Conservative
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Matches 8; Conservative
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US-10-258-144-317
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; Sequence 297, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Livingston, Phil
; APPLICANT: Mayhew, Mark
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA; TITLE OF INVENTION: BROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA; TITLE OF INVENTION: BROTE: 2002-10-17
; PRIOR APPLICATION NUMBER: 60/197,642
; PRIOR PILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: FRAESEQ for Windows Version 3.0
; SEQ ID NO 297
; LENGTH: 17
; TYPE: DATE: 2001-04-17
; TYPE: DATE: 2001-04-17
; TYPE: DATE: 2001-04-17
; TYPE: DATE: 2001-04-17
; TYPE: DATE: 2001-04-17
; TYPE: DATE: 2001-04-17
                                                                                                         RESULT 42
US-10-258-144-288
; Sequence 288, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
   APPLICANT: Al-Awgati, Qais
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Maybew, Mark
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Maybew, Mark
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
; TITLE OF INVENTION: PROTEIN MEDIATE INVENTION: PROTEIN MEDIATE 2002-10-17
; CURRENT PRING DATE: 2002-10-17
; PRIOR PILING DATE: 2000-04-17
; PRIOR FILING DATE: 2000-04-17
; SEQ ID NO 288
; LEASTHARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 288
; LYPE: PRT
; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-288
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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US-10-258-144-297
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US-10-258-144-298
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
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Best Local Similarity 100.
Matches 8; Conservative
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Pain

APPLICANT: Houghton, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: How, Mark

APPLICANT: How, Mark

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: PROTEIN INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR PRILING DATE: 2000-04-17

PRIOR PILING DATE: 2000-04-17

PRIOR PILING DATE: 2000-04-17

SEQ ID NO 327

LENGTH: 17
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APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Meo, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERBERS: 11746/4401
CURRENT APPLICATION NUMBER: 105/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FASTSEQ FOR Mindows Version 3.0
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mee, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 64; DB 4; Length 17; Best Local Similarity 100.0%; Pred. No. 0.062;
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                  SEQ ID NO 318
| LENGTH: 17
| TYPE: PRT
| ORGANISM: Homo Bapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT; ORGANISM: Homo sapiens derivative US-10-258-144-327
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-328
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US-10-258-144-388

| Sequence 389, Application US/10258144
| Publication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Al-Awqati, Qais
| APPLICANT: Maybew, Mark
| APPLICANT: Maybew, Mark
| TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOWA
| TITLE OF INVENTION: PROTEIN WEDIATED INMUNOTHERAPY OF MELANOWA
| FILE REFERENCE: 11146/46401
| CURRENT FILING DATE: 2002-10-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LIVINGEON, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hos, Mark
APPLICANT: Hos, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA;
FILER REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PASESEQ for Windows Version 3.0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 387, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
                                                                                                                                                                                                                       LENGTH: 17
TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-328
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Gaps

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RESULT 54

US-10-258-144-428

i Sequence 428, Application US/10258144

i Publication No. US20040101532A1

i GENERAL INFORMATION:

APPLICANT: Houghton, Alan

i APPLICANT: Houghton, Phil

APPLICANT: How Mark

APPLICANT: How Mark

i APPLICANT: How Mark

i APPLICANT: Al-Awdati, Qais

APPLICANT: How Mark

i TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

i TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

i FILE REFERENCE: 11746/46401

cURRENT APPLICATION NUMBER: US/10/258,144

cURRENT FILING DATE: 2000-10-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

i NUMBER OF SEQ ID NOS: 502

i SOFTWARE: FastSEQ for Windows Version 3.0

i FONTWARE: PastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                     Sequence 427, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: NAYBERIN BEOTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
FILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502

SOFTHARE: FRELENCE FRELENCE OF Windows Version 3.0
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                                                                         100.0%; Score 64; DB 4; Length 17;
100.0%; Pred. No. 0.062;
/ative 0; Mismatches 0; Indels
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; ORGANISM: Homo sapiens derivative US-10-258-144-418
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; ORGANISM: Homo sapiens derivative
US-10-258-144-428
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                                                                                                 Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
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Best Local Similarity
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US-10-258-144-427
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Publication No. US20040101532A1

GENERAL INPORMATION:

APPLICANT: LIVINGROON, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: MayNew, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF ILVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF ILVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF ILVENTION: METHODS AND COMPOSITIONS FOR MELANOMA

FILE REPRENCE: 11146/46401

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FREUSED FOR WINGOWS VERSION 3.0

SEQ ID NO 417

LENGTH: 17
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Howardti, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46101
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 418
LENGTH: 17
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100.0%; Score 64; DB 4; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0; Indels
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; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: PastSEQ for Windows Version 3.0
; SEQ ID NO 388
; LENGTH: 17
; TYPE: PRT
TYPE: PRT
US-10-258-144-388
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Best Local Similarity 100.º
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US-10-258-144-418
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10 HWDFAWPW 17
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  Gaps
                                                                                                                                                                                                                                 APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awdati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Most Mark
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46101
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-04-17
PRIOR PLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 437
LENGTH: 17
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APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Houghton, Pair
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: How, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/358,144
CURRENT FILING DATE: 2002-10-17
PRIOR PAPLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: FREESEQ for Windows Version 3.0
SEQ ID NO 438
LENGTH: 17
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  Indels
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Mismatches
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Publication No. US20040101532A1
GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens derivative
US-10-258-144-437
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Best Local Similarity luv...
8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
8; Conservative
                                                             10 HWDFAWPW 17
                                     1 HWDFAWPW 8
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Matches
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10 HWDFAWPW 17

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                                                                         | APPLICANT: Houghton, Alan |
| APPLICANT: Houghton, Phil |
| APPLICANT: Livingston, Phil |
| APPLICANT: Al-Aqadi, Qais |
| APPLICANT: Al-Aqadi, Qais |
| APPLICANT: Al-Aqadi, Qais |
| APPLICANT: Mayhew, Mark |
| APPLICANT: Hoe, Mee |
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK |
| TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA |
| TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA |
| TITLE OF INVENTION: PROTEIN UNDBER: US/10/258,144 |
| CURRENT APPLICATION NUMBER: 60/197,642 |
| PRIOR PILING DATE: 2000-04-17 |
| NUMBER OF SEQ ID NOS: 502 |
| SEQ ID NO 467 |
| LENGTH: 17 |
| TYPE: PRI |
| ORGANISM: Homo sapiens derivative |
| US-10-258-144-467 |
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
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Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 8; Conservative 0; Mismatches 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 468
LENGTH: 17
Sequence 467, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
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; Sequence 497, Application US/10258144
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Best Local Similarity 100.
Matches 8; Conservative
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Gaps

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APPLICANT: HOLLING, METHODS AND COMPOSITIONS FOR HEAT SHOCK TITLE OF INVENTION: PREHEDS AND COMPOSITIONS FOR HEAT SHOCK TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PRAESEQ for Windows Version 3.0
SEQ ID NO 267
LENGTH: 18
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Publication No. US20040101532A1
Publication No. US20040101532A1
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Aydew, Mark
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/446101
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.065;
/ative 0; Mismatches 0; Indels
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APPLICANT: Mayhew, Mark
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US-10-258-144-307
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Sequence 498, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

CURENT: Mayhew, Mark

CURRENT PILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2000-04-17

PRIOR PILING DATE: 2000-04-17

NUMBER: FOOL DATE: 2000-04-17

SEQ ID NO 498

LENGTHARE: FOOLESE FOOLESE

LENGTH: 17

MARKET FOOLESE

LENGTH: 17

MARKET FOOLESE

SEQ ID NO 498
         GENERAL INFORMATION:
APPLICANT: Houghcon, Alan
APPLICANT: Houghcon, Alan
APPLICANT: Houghcon, Alan
APPLICANT: Al-Awgatio, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11146/46401
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2000-40-17
NUMBER OF SEQ ID NOS: 502
SOCTION OF SEQ ID NOS: 502
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SOCTION OF SEQ ID NOS: 502
SEQ ID NO 497
LENGTH: 17
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100.0%; Pred. No. 0.062;
tive 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.062;
/ative 0; Mismatches 0; Indels
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-497
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Publication No. US20040101532A1
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-267
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Me
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-44-17
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR TILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-04-17
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 338
LENGTH: 18
TYPE: PRT
ORGANISW: Homo sapiens derivative
US-10-258-144-338
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 347
LENGTH: 18
          SOFTWARE: FastSEQ for Windows Version 3.0
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Publication No. US20040101532A1
GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
                               ; SEQ ID NO 337
; LENGTH: 18
; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-337
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ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Marhaw, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT ELLING DATE: 2002-10-17
CURRENT ELLING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PELICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 308
LENGTH: 18
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Sequence 337, Application US/10258144

Publication No. US20040101532A1

Sequence 337, Application US/10258144

Publication No. US20040101532A1

SENERAL INFORMATION:

APPLICANT: Livingston, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Hoe, Me

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17

PRIOR PILICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PSESSEQ for Windows Version 3.0
SOFTWARE: PSESSEQ for Windows Version 3.0
IENGTH: 18
TYPE: PRT
TYPE: PRT
US-10-258-144-307
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; ORGANISM: Homo sapiens derivative
US-10-258-144-308
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Best Local Similarity luv...
8, Conservative
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Gaps

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APPLICANT: Livingston, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPRENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 367
                                                                                                                                                                                                                                                                                                                                                                                                                     PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
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                                                                                                                                                                              Sequence 358, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton,

APPLICANT: Houghton,

APPLICANT: Houghton,

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

FILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOC

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT

CURRENT FILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT

CURRENT FILLE OF INVENTION NUMBER: US/10/258,144

CURRENT FILLING DATE: 2000-04-17

PRIOR PELING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SEQ ID NO 358

SEQ ID NO 358

LENGTH: 18
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Publication No. US20040101532A1
GENERAL INFORMATION:
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Best Local Similarity 100.
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                       HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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| Deblication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Livingston, Phil
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOWA
| TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOWA
| TITLE OF INVENTION NUMBER: US/10/258,144
| CURRENT APPLICATION NUMBER: 60/197,642
| PRIOR FILING DATE: 2000-04-17
| NUMBER OF SEQ ID NOS: 502
| SEQ ID NO 357
| LENGTH: 18
| TYPE: PRI
| TYPE: PRI
| CRGANISM: Homo sapiens derivative
| US-10-258-144-357
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ILVINGBLOM, ALGERIA, APPLICANT: LIVINGBLOM, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: PRACEED FOR Windows Version 3.0
SEG ID NO 348
LENGTH: 18
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                                              100.0%; Score 64; DB 4; Length 18; 100.0%; Pred. No. 0.065;
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                                                                                                0; Indels
                                                                                                  0; Mismatches
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; ORGANISM: Homo sapiens derivative
US-10-258-144-348
                                                                        Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-347
                                                 Query Match
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: MUMBER: 0100/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500-04-17
NUMBER OF SEQ ID NOS: 500-04-17
SOFTWARE: PSECSEQ FOR Windows Version 3.0
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Publication No. US20040101532A1

GENERAL INFORMATION,

APPLICANT: Houghton, Phil

APPLICANT: Livingston, Phil

APPLICANT: Al-awqari, Qais

APPLICANT: Maybew, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT PRING DATE: 2000-10-17

PRIOR FILING DATE: 2000-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 64; DB 4; Length 18; Best Local Similarity 100.0%; Pred. No. 0.065; Matches 8; Conservative 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 397
LENGTH: 18
TYPE: PRT
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 18
TYPE: PRT
ORGANISM: Homo sapiens derivative
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Al-Awgati, Qais
Mayhew, Mark
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APPLICANT: Livingston, Ph
APPLICANT: Al-Awgati, Qai
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                                                                                                                                                         APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: MayNew, Mark
APPLICANT: MayNew, Mark
APPLICANT: MayNew, Mark
APPLICANT: MayNew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: 1746,4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/27,642
PRIOR PILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 368
LENGTH: 18
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Publication No. US20040101532A1
GENERAL INFORMATION;
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagati, Qais
APPLICANT: Al-Awagati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: MORBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 377
TYPE: PRT
TYPE: PRT
TYPE: PRT
TYPE: PRT
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; Sequence 378, Application US/10258144
; Publication No. US20040101532A1
                                                       Sequence 368, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ·TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-368
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
Matches 8; Conserv
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; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-408
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Best Local Similarity 100..
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Best Local Similarity 100.0
Matches 8; Conservative
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US-10-258-144-448
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Al-Awqati, Qais

APPLICANT: Maybew, Mark

TITLE OF INVENTION: BRITHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT FILING DATE: 2002-10-17

FRIOR PILLING DATE: 2002-10-17

FRIOR PILLING DATE: 2002-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PRESENCE: 2000-04-17

SEQ ID NO 407

LENGTH: 18
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APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PARKEQ for Windows Version 3.0
SEQ ID NO 398
LENGTH: 18
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
APPLICANT: Hoe, Mer
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APPLICANT: Houghton, Phil
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 9; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens derivative US-10-258-144-398
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; ORGANISM: Homo sapiens derivative
US-10-258-144-407
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Best Local Similarity 100.
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US-10-258-144-408
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1 Sequence 447, Application US/10258144

1 Sequence 447, Application US/10258144

1 Publication No. US/20040101532A1

1 GENERAL INFORMATION:

1 APPLICANT: Houghton, Alan

1 APPLICANT: Al-Awqati, Qais

2 APPLICANT: Al-Awqati, Qais

3 APPLICANT: Al-Awqati, Qais

3 APPLICANT: Al-Awqati, Qais

4 APPLICANT: Mayhew, Mark

5 TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

7 TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

7 TITLE OF INVENTION: PROTEIN WIMBER: US/10/258,144

7 CURRENT PILING DATE: 2000-210-17

7 PRIOR APPLICATION NUMBER: 60/197,642

7 PRIOR PILING DATE: 2000-04-17

7 NUMBER OF SEQ ID NOS: 502

7 SEQ ID NOS: 502

7 SEQ ID NOS: 502

7 SEQ ID NOS: 502

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7 SEQ ID NOS: 502

7 SEQ ID NOS: 502
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Houghton, Phil

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Hoe, Wee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 408
LENGTH: 18
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Gaps

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Sequence 478, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Phil

APPLICANT: Houghton, Phil

APPLICANT: Houghton, Phil

APPLICANT: How, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOWA

FILE REPRENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT PELICATION NUMBER: 60197,642

PRIOR FILING DATE: 2000-04-17
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERBLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-24-17
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
                       100.0%; Score 64; DB 4; Length 18; 100.0%; Pred. No. 0.065;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 478
LENGTH: 18
                                                                                                                                                                                                                                                                                                    Sequence 477, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 18
TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Homo sapiens derivative
                       Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                          11 HWDFAWPW 18
                                                                                                                            1 HWDFAWPW 8
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Publication No. US20040101532A1

GENERAL INFORMATION;
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTOTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLILING DATE: 2000-04-17
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: How, Mark
APPLICANT: How, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FLIE REPERBURE: 1746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 458
LENGTH: 18
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                                          TYPE: PRT ; ORGANISM: Homo sapiens derivative US-10-258-144-448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens derivative US-10-258-144-457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-458
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                                                                                                                                                                                                                                                    1 HWDFAWPW 8
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SEQ ID NO 448
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LENGTH: 18
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RESULT 0.
US-09-794-517-8

| Sequence 8, Application US/09794517
| Publication No. US20030021794A1
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| HAPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E.
| HARTL, F. Ulrich HOUGHPON, Alan HOUGHPON, Alan TAKECHI, Yoshizumi
| HOUGHPON, Alan TAKECHI, Yoshizumi
| HOUGHPON, Mark HOAR Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOCTWARRE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: AUGUST: 60/002,490
FILING DATE: AUGUST: 18, 1995
APPLICATION NUMBER: 09/011,645
FILING DATE: AUGUST: 18, 1995
ATTORNEY/AGENT INCPRANTION:
NAME: DeLUCIA: Richard L.
REGISTRATION NUMBER: 28, 839
FELECOMMUNICATION NUMBER: 28, 839
FELECOMMUNICATION NUMBER: 28, 839
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 
TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 3; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
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US-09-794-517-9
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| US-10-289-144-48B
| Publication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Livingston, Phil
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| APPLICANT: Maynew, Mark
| TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
| TITLE OF INVENTION: PROTEIN WEBBR: US/10/258,144
| CURRENT PEPLICATION NUMBER: 60/197,642
| PRIOR FILING DATE: 2000-04-17
| NUMBER OF SEQ ID NOS: 502
| SOFTWARE: FastSEQ for Windows Version 3.0
| SEQ ID NO 488
| LYPE: PRT | TYPE: PRT |
| ORGANISM: Homo sapiens derivative
| US-10-258-144-488
                                                                                                                                                                                                                                                                                                                          APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: (0/197,642)
PRIOR PILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 487
LENGTH: 18
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100.0%; Score 64; DB 4; Length 18;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                      , Sequence 487, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT; ORGANISM: Homo sapiens derivative US-10-258-144-487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                         HWDFAWPW 18
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11 HWDFAWPW 18
HWDPAWPW 8
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
COMPUTER: SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIFICATION ATA:
REOR APPLICATION DATA:
REOR APPLICATION DATA:
APPLICATION UWBER: US/09/011,645
                                                                                                        HOUGHTON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotheraples
                     Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 64; DB 3; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
RADDRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 9, Application US/09794529
; Publication No. US20030082197A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION:
                                                                                       HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 10004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 HWDFAWPW 19
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      GENERAL INFORMATION:
                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-794-529-9
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                                                                                                                                                                                                            TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
Sequence 9, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HARTL, F. Ulrich
HOE, Mee H.
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 3; Length 19; 100.0%; Pred. No. 0.068;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: «UDKNOWN»
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: AUGUST 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEFAK: (212) 425-7200
TELEFAX: (212) 425-528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA: APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 89
US-09-794-529-8
Sequence 8, Application US/09794529
Publication No. US20030082197A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: <Unknown>
                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                           CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-794-517-9
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ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                      APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 64; DB 3; Length 19; 100.0%; Pred. No. 0.068; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM compatible
CORREATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 28, 839
REGISTRATION NUMBER: 28, 839
REGISTRATION NUMBER: 28, 839
REGISTRATION NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212) 425-720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 19
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Publication No. US20030082198A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8; Conservative
                                                                                                                                                                                                                                                                                                           CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 HWDFAWPW 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW 8
  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 92
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               APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ultich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.0%; Score 64; DB 3; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0. Tradala
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
PILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM compatible
OPERATING SYSTEMS, MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 8, Application US/09794832 ; Publication No. US20030082198A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 425-5288
TELEX: «UNKNOM»
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                           CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 HWDFAWPW 8
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GENERAL INFORMATION:
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US-09-794-832-8
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Gaps

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APPLICANT: HOUGHOON, Alan
APPLICANT: HOUGHOON, Alan
APPLICANT: HOUGHOON, Alan
APPLICANT: HOUGHOON, Alan
APPLICANT: Out-fell 10 Out-hek
TYTLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
FILE REFERENCE: 11746/46004
CURRENT APPLICATION NUMBER: US/10/053,520
CURRENT FILING DATE: 2002-10-01
PRIOR PILING DATE: 1997-10-31
NUMBER OF SEQ ID NOS: 321
SOFFWARE: WordPerfect 8.0 for Windows
SEQ ID NO 321
LENGTH: 19
                                                                    APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Hartl, Ulrich
APPLICANT: Ouerfelli, Ouathek
APPLICANT: Moroi, Yoichi
TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 64; DB 4; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
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                   loan-Kettering Institute for Cancer Research Rothman, James E. Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sloan-Kettering Institute for Cancer Research APPLICANT: Rothman, James E. APPLICANT: Mayhew, Mark
                                                                                                                                                                                                                             FILE REFERENCE: 11746/4603
CURRENT APPLICATION NUMBER: US/10/052,578
CURRENT FILING DATE: 2002-01-17
FRIOR APPLICATION NUMBER: 08/961,707
PRIOR FILING DATE: 1997-10-31
NUMBER OF SEQ ID NOS: 321
SOFTWARE: WordPerfect 8.0 for Windows
; SOFTWARE: 1997-110-31
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S-10-053-520-321
Sequence 321, Application US/10053520
Publication No. US20030166530A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
; OTHER INFORMATION: hybrid peptide
US-10-052-578-321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
; OTHER INFORMATION: hybrid peptide
US-10-053-520-321
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 8; Conservative
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12 HWDFAWPW 19
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                 APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, P. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 3; Length 19;
llarity 100.0%; Pred. No. 0.068;
Conservative 0; Mismatches 0: Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb.1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
8; Conserve
GENERAL INFORMATION:
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Gaps

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Gaps

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RESULT 95

US-10-052-578-321 ; Sequence 321, Application US/10052578 ; Publication No. US20030134787A1

RESULT 93

HWDFAWPW

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FEATURE
  JS-10-170-713A-9
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               Sequence 8, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
APPLICANT: GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 19;
100.0%; Pred. No. 0.068;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: CABBON, LAWRENCE P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4

COMPUTER: IBM compatible

OPERATING SYSTEM: MS DOS

SOFTWARE: Word Perfect

CURRENT APPLICATION DATA:

PILLING DATE: 13-40n-2002

CLASSIFICATION: A24

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
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SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION
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                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 30
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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Best Local Similarity 100.
Matches 8; Conservative
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JS-10-170-713A-8
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Sequence 9, Application US/1017013A
Publication No. US20030185842A1
GENERAL INFORMATION:
APPLICANT: SLOAN-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKEGHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 19; 100.0%; Pred. No. 0.068; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Casson, Lawrence P.
REGISTATION NUMBER: 46,606
REPERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
PILING DATE: 13-7un-2002
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
PELING DATE: February 13, 1998
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
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                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 30
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                       CITY: New York
STATE: NY
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HWDFAWPW 19

RESULT 96

RESULT 97

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HARTL, F. Ulrich
HOB, Mee H.
HOGHTON, Alan
TAKEHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                              Sequence 9, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 64; DB 4; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/10/171,734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS
LENGTH: 19
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
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                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                               CITY: New York
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                            US-10-171-734-9
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Sequence 8, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOK, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MATHEW, MARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 64; DB 4; Length 19; 100.0%; Pred. No. 0.068;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM COMPACTURED COMPUTER: IBM COMPACTURED COMPACTURED COURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
PPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
                                                                                                                                                                                                                                              Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESSS:
RADDRESSRES Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lawrence P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL: no
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                  SEOUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Casson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |||||||||
HWDFAWPW 19
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                           STATE: NY
                                                                                                                                                                                                                                                                  NUMBER OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-171-734-8
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US-10-258-147-17
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                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 64; DB 4; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                          US-10-053-498B-321
Sequence 321, Application US/10053498B
Sequence 321, Application No. US20030194409A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-258-147-16

Sequence 16, Application US/10258147

Publication No. US20040043419A1

GENERAL INPORMATION:

APPLICANT: Rochman, James

APPLICANT: Hoe, Mee

TITLE OF INVENTION: JAVELINIZATION OF PROTEINS

FILE REFERENCE: 11746/46276

CURRENT PILING DATE: 2002-10-17

CRRIOR FILING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 47

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 16

LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: hybrid peptide US-10-053-498B-321
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; OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin
US-10-258-147-17
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100.0%; Score 64; DB 4; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
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Sequence 17, Application US/10258147

Publication No. US20040043419A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: ROthman, James
APPLICANT: Hoe, Me
ITILE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE OF INVENTION JAVELINIZATION OF PROTEINS
FILE OF INVENTION UNMBER: US/10/259,147

CURRENT FILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR APPLICATION NUMBER: 60/197,462

RADOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 47

SEQ ID NO 17

LENGTH: 19
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Sequence 22, Application US/10258147
Publication No. US20040043419A1
GENERAL INFORMATION
TITLE GANT: Rothman, James
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT PILING DATE: 2002-10-17
CURRENT FILING DATE: 2002-10-17
FRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: FaetSEQ for Windows Version 3.0
SEQ ID NO 22
LENGTH: 19
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Publication No. US20040043419A1
GENERAL INFORMATION:
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ORGANISM: Aritifical Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich

APPLICANT: Hood, Mee H.

APPLICANT: Houghton, Alan

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461061

CURRENT APPLICATION NUMBER: US/10/367,580
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US-10-258-147-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 64; DB 4; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
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                   APPLICANT: Hos, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION UNWBER: US/10/258,147
CURRENT PILING DATE: 2002-10-17
PRIOR PELLING DATE: 2000-04-17
PRIOR PELLING DATE: 2000-04-17
NUMBER: FeetSEQ for Windows Version 3.0
SOFTWARE: FeetSEQ for Windows Version 3.0
IENGTH: 19
TYPE: PRI
ORGANISM: Aritifical Sequence
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PRIOR APPLICATION NUMBER: US 09/794,832

PRIOR FILING DATE: 2001-02-7

PRIOR FILING DATE: 1998-02-13

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR PILING DATE: 1998-02-13

PRIOR PILING DATE: 1996-08-16

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

SEQ ID NO 8

LENGTHALE: WordPerfect 8.0 for Windows

SEQ ID NO 8

LENGTH: 19
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ORGANISM: Artificial Sequence
FEATURE:
Mayhew, Mark
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ઠે 셤 RESULT 105 US-10-367-580-9

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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Hoebthon, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
ITILE OF INVENTION: Hoat Shock Protein-Based Vaccines and Immunotherapies
ITILE OF INVENTION: Hoat Shock Protein-Based Vaccines and Immunotherapies
ITILE OF INVENTION: Hoat Shock Protein-Based Vaccines and Immunotherapies
FIER REFERENCE: 1746/461012
CURRENT APPLICATION NUMBER: US 00/01,645
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
SPRIOR PILING DATE: 1995-08-18
APPLICANT: Rothman, James E.
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mark
TITLE OF INVENTION: Hat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461061
CURRENT APPLICATION NUMBER: US/10/367,580
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,832
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1995-08-16
PRIOR PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR APPLICATION NUMBER: US 60/002,479
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; OTHER INFORMATION: synthetic peptide
US-10-367-580-9
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OTHER INFORMATION: synthetic peptide
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 8: Conserv
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Squence 9, Application US/10367593

Publication No. US20040071721A1

GENERAL INFORMATION:

APPLICANT: Rothman, James B.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Mayhew, Mark

ITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REPERBNCE: 11746/461012

CURRENT APPLICATION NUMBER: US/10/367,593

CURRENT APPLICATION NUMBER: US 09/011,645

PRIOR PILING DATE: 1996-02-13

PRIOR PILING DATE: 1996-08-16

PRIOR PILING DATE: 1995-08-16

SOFTWARE: WordPerfect 8.0 for Windows

SEQ ID NO 9

LENGTH: 199
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Publication No. US20040071722A1

GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Haybew, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461041

CURRENT APPLICATION NUMBER: US/10/367,594

CURRENT APPLICATION NUMBER: US/2004680,806

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 1998-02-13

PRIOR PLICATION NUMBER: PCT/US96/13363
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                                       100.0%; Score 64; DB 4; Length 19; 100.0%; Pred. No. 0.068; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                           Local Similarity 100.
1es 8; Conservative
                                                                                                                                                       12 HWDFAWPW 19
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US-10-367-593-9
US-10-367-593-8
                                         Query Match
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Matches
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APPLICANT: Rothman, James E.
APPLICANT: Rothman, James E.
APPLICANT: Howelve, Mee H.
APPLICANT: Howelve, Mea H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark
ITILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461041
CURRENT APPLICATION NUMBER: US/10/367,594
CURRENT APPLICATION NUMBER: US 09/680, 806
PRIOR PILING DATE: 2000-10-05
PRIOR PILING DATE: 1998-02-13
PRIOR PRILING DATE: 1996-08-16
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
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PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
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100.0%; Pred. No. 0.068;
vative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
PRIOR FILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: synthetic peptide US-10-367-594-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTHER INFORMATION: synthetic peptide US-10-367-594-9
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US-10-367-654-8
Sequence 8, Application US/10367654
Publication No. US20040071723A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9, Application US/10367594 Publication No. US20040071722A1 GENERAL INFORMATION:
APPLICANT: Rothman, James E.
                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Sequence 8, Application US/10367658 Publication No. US20040071724A1 GENERAL INFORMATION:
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Publication No. US20040071724A1
GENERAL INFORMATION:
    ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hath, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US/10/367,654
FILE REFERENCE: 10746/461032
CURRENT FILING DATE: 2002-06-13
FRIOR APPLICATION NUMBER: US 09/613,295
FRIOR PILING DATE: 1996-08-16
FRIOR PILING DATE: 1996-08-16
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
IIIE REFERENCE: 11746/461032
CURRENT APPLICATION WUMBER: US/10/367,654
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION WUMBER: US 10/171,734
PRIOR PILING DATE: 2000-06-13
PRIOR PILING DATE: 2000-08-10
PRIOR APPLICATION WUMBER: US 09/0316,295
PRIOR APPLICATION WUMBER: US 09/031,645
PRIOR PILING DATE: 1998-02-16
PRIOR PILING DATE: 1998-02-16
PRIOR APPLICATION WUMBER: US 60/002,490
PRIOR APPLICATION WUMBER: US 60/002,490
PRIOR APPLICATION WUMBER: US 60/002,479
PRIOR APPLICATION WUMBER: US 60/002,479
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
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100.0%; Pred. No. 0.068;
vative 0; Mismatches 0; Indels
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; OTHER INFORMATION: synthetic peptide
US-10-367-654-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
Rothman, James E.
Hartl, F. Ulrich
Hoe, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION NUMBER: US/10/367,658
CURRENT APPLICATION NUMBER: US 09/794,529
PRIOR APPLICATION NUMBER: US 09/794,529
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR PRILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
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APPLICANT: Rath, V. Ulrich
APPLICANT: Harl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mar.
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REPERENCE: 1746/461051
CURRENT APPLICATION NUMBER: 2003.02-14
CURRENT FILING DATE: 2003.02-14
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                                                                                        Query Match 100.0%; Score 64; DB 4; Length 19; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
; OTHER INFORMATION: synthetic peptide US-10-367-654-9
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SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 8
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US-10-367-658-8
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US-10-258-144-274
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publication No. US20040071725A1

GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, James E.
APPLICANT: Hard, Yoshizum
APPLICANT: Houghton, Yoshizum
APPLICANT: Mayhaw, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461072
CURRENT APPLICATION NUMBER: US/10/367,668
CURRENT PILING DATE: 2003-02-13
FRIOR APPLICATION NUMBER: US 09/794,517
FRIOR PILING DATE: 1996-02-13
FRIOR PILING DATE: 1996-02-13
FRIOR PILING DATE: 1996-08-16
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100.0%; Score 64; DB 4; Length 19;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 8; Conservative 0; Mismatches 0; Indels
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                PRIOR AFFLING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: PCT/US96/1363
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
LENGTH: 19
APPLICATION NUMBER: US 09/794,529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTHER INFORMATION: synthetic peptide US-10-367-658-9
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STATEMENT IN CORTAIN OF A PROPERTY OF A PRESENCE IN CONTRACTOR IN CORTAIN OF A PRESENCE IN CONTRACT Hartl, F. Ulrich

A APPLICANT: Hartl, F. Ulrich

A APPLICANT: Houghton, Alan

A APPLICANT: Houghton, Alan

A APPLICANT: Houghton, Alan

A APPLICANT: Mark

ITILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461021

CURRENT APPLICATION NUMBER: US/10/367,674

CURRENT PILING DATE: 2003-06-13

PRIOR APPLICATION NUMBER: US 09/55,868

PRIOR APPLICATION NUMBER: US 09/51,868

PRIOR PILING DATE: 1996-02-13

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR APPLICATION NUMBER: US 06/002,490

PRIOR FILING DATE: 1995-08-16

PRIOR FILING DATE: 1995-08-18

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR FILING DATE: 1995-08-18

PRIOR APPLICATION NUMBER: US 60/002,479
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                                                                                                                                                                                                                                                                                                      100.0%; Score 64; DB 4; Length 19; 100.0%; Pred. No. 0.068; tive 0; Mismatches 0; Indels
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Publication No. US20050202033A1
GENERAL INFORMATION:
APPLICANT: Fletchmer, Jessica
APPLICANT: Prince-Cohane, Kenya
     PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 8
                                                                                                                                                                                             FRATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-674-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/10367674 Publication No. US20040127684A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 HWDFAWPW 19
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US-10-776-521B-368
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FILE REFERENCE: 11746/4610211 (2/10/367,674)

FILE REFERENCE: 11746/4610211 (2/10/367,674)

FRIOR APPLICATION NUMBER: US 10/170,738

FRIOR APPLICATION NUMBER: US 09/011,645

FRIOR PRILING DATE: 2000-04-20

FRIOR FILING DATE: 1998-02-13

FRIOR FILING DATE: 1996-08-16

FRIOR APPLICATION NUMBER: US 60/002,490

FRIOR APPLICATION NUMBER: US 60/002,490

FRIOR FILING DATE: 1995-08-18

FRIOR FILING DATE: 1995-08-18
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Meyhew, Mark
APPLICANT: Meyhew, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT PLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-4-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 275
LENGTH: 19
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100.0%; Score 64; DB 4; Length 19; larity 100.0%; Pred. No. 0.068; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; LOCATION: (9)...(11)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-275
                                                                                                                                                                                                                                                                         5-10-258-144-275
Sequence 275, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8, Application US/10367674 Publication No. US20040127684A1 GENERAL INFORMATION:
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                            Best Local Similarity
Matches 8; Conser
                                                                                                                                                1 HWDFAWPW
     Query Match
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Gaps

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ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                          US-10-776-521B-379
Sequence 379, Application US/10776521B
; Publication No. US20050202033A1
; GENERAL INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Hybrid antigen US-10-776-521B-379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                         12 HWDPAWPW 19
                    HWDFAWPW 8
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### APPLICANT: Mente, suil.
### APPLICANT: Madjele, Sofija
### APPLICANT: Slusarewicz, Paul
#### APPLICANT: Barber, Brian
#### APPLICANT: Barber, Brian
#### TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
### TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
### TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
### TITLE OF INVENTION: IMPROPER: 2004-02-12
### CURRENT APPLICATION NUMBER: 60/503,417
### PRIOR PLING DATE: 2003-09-16
### PRIOR PLING DATE: 2003-09-16
### PRIOR PLING DATE: 2003-04-18
### PRIOR PLING DATE: 2003-04-18
### PRIOR PLING DATE: 2003-04-13
### PRIOR PLING DATE: 2003-02-13
### NUMBER OF SEQ ID NOS: 419
### SOFTWARE: PastSEQ for Windows Version 4.0
### ILENGTH: 19
### ILENGTH: 19
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US-10-776-521B-376
Sequence 376, Application US/10776521B
Sequence 376, Application US/10776521B
Sequence 376, Application No. US20050202033A1
GENERAL INFORMATION:
APPLICANT: Fletchner, Jessica
APPLICANT: Prince-Chane, Kenya
APPLICANT: Andielic, Souil
APPLICANT: Briden Solida
APPLICANT: Barber, Solida
TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF LIVENTION: IMMUNOTHERAPIES
TITLE OF LIVENTION: IMMUNOTHERAPIES
TITLE OF LIVENTION: UNMUNER: US/10/776,521B
CURRENT FILING DATE: 2004-02-12
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PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-04-18
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-06-13
SOFTWARE: PASESEQ for Windows Version 4.0
SEQ ID NO 376
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Hybrid antigen US-10-776-521B-368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , OTHER INFORMATION: Hybrid antigen US-10-776-521B-376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 HWDFAWPW 19
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GENERAL INVOKATION:

GENERAL INVOKATION:

APPLICANT: Fletchner, Jessica
APPLICANT: Prince-Cohane, Kenya
APPLICANT: Prince-Cohane, Kenya
APPLICANT: Mehra, Sunil
APPLICANT: Madjelic, Sofija
APPLICANT: Andjelic, Sofija
APPLICANT: Baber, Balan
ITILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
ITILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES
CURRENT APPLICATION NUMBER: US/10/76,521B
CURRENT APPLICATION NUMBER: US/10/76,521B
CURRENT APPLICATION NUMBER: 60/463,417
PRIOR FILING DATE: 2003-09-16
PRIOR PILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR SEQ ID NOS: 419
SOFTWARE: PASCEQ for Windows Version 4.0

SEQ ID NO 379
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US-09-744-517-10
| Sequence 10, Application US/09794517
| Publication No. US20030021794A1
| Publication No. US20030021794A1
| GENERAL INFORMATION:
| APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich
| HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi MAYREW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOUSTHON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 12, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
ROTHWAN, James B.
HOREL, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPES. Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
       SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
CLASSIFICATION DATA:
RAPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: <UNKNOWN:
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virus vaccine
;
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
US-09-794-517-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDENDESS: CURNOWN>
TOPOLLGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                       FELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-794-517-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HATLL, F. Ulrich
HOE, Mee H.
HOKE, Mee H.
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 64; DB 3; Length 20; ilarity 100.0%; Pred. No. 0.071; Conservative 0; Mismatches 0; Indels
                                                                   virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                     FELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-794-517-11
; Sequence 11, Application US/09794517
; Publication No. US20030021794A1
; GENERAL INFORMATION:
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                       FELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECILE TYPE: peptide
HYPOTHETICAL: yes
FRAGWENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
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OTHER INFORMATION: hybrid peptide for human papilloma
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Publication No. US20030021794A1
GENERAL INPORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEN, Mark
TITLE OF INVENTION: Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                CURKENT APPLICATION DATA:

PULING DATE: 27-Feb-2001
CLASSIFICATION CUNKNOWN>
PILING DATE: 27-Feb-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: cUnknown>
APPLICATION NUMBER: 01995
APPLICATION NUMBER: 13996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEFONE: (212) 425-7200
TELEFONE: CUNKNOWN>
INFORMATION FOR SEQ ID NO: 13:
SEGUBENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  yirus vaccine
;
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-794-517-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-794-517-14
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Publication No. US20030021794A1

GENERAL INFORMATION:

HAPPLICANT: Sloan *Kettering Institute for Cancer Research
ROTHMAN, James E.

HARTL, F. Ulrich
HOE, Mee H.

HOUGHTON, Alan

TAKECHI, Yoshizumi
MAYHEW, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies

Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
ZIP: 10004
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
CORRUTER: IBM compactible
COMPUTER: IBM compactible
CORRATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 64; DB 3; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                          PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: 09/011,645
FILING DATE: AUGUST: 09/011,645
FILING DATE: AUGUST: 60/002,490
FILING DATE: AUGUST: 18, 1995
APPLICATION NUMBER: 60/1996
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEPHONE: (212) 425-5288
TELEFRA: CURRONN:
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27.Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus vaccine

sequence Description: Seq ID No: 12:
US-09-794-517-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: < Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 30
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13 HWDPAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE
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0; Gaps

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OTHER INFORMATION: hybrid peptide for human papilloma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James E.
HARTL, F. Ulrich
HORE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIF: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071; tive 0; Mismatches 0; Indels
                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: AUGNOWN>
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus vaccine, sequence Description: SEQ ID NO: 15: US-09-794-517-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECTLE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                 FELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                 TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWDFAWPW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JS-09-794-517-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN. James B.
HARTL, F. Ulrich
HOB, Mee H.
HOGHTON, Alan
TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                    FILING DATE: «Unknown)
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLAGIA, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
APPLICATION NUMBER: US/09/794,517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                    FILING DATE: 27-Feb-2001
CLASSIPTCATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
RADBRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette - COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virus vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |||||||||
HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-794-517-14
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RESULT 131
US-09-745-17-18
| Sequence 18, Application US/09794517
| Publication No. US20030021794A1
| Publication No. US20030021794A1
| GENERAL INFORMATION:
| APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich
| HOR, Mee H. HOUGHPON, Alan TAKECHI, Yoshizumi MATHEW, Mark Mark MATHEW, Mark Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 64; DB 3; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 8; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/011,645
FILING DATE: «Unknown»
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: «Unknown»
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILLING DATE: 27-Feb-2001
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
sequence description: Seq ID NO: 17:
US-09-794-517-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: NY
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US-09-794-517-17
Sequence 17, Application US/09794517
Sequence 17, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,517
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                  FILING DATE: «UNKNOWN)

PILING DATE: «UNKNOWN)

APPLICATION NUMBER: 60/002,490

FILING DATE: August 18, 1995

APPLICATION NUMBER: PCT/USS6/13363

FILING DATE: August 16, 1996

ATTORNEY/ABGNT INFORMATION:
NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839

REFRENCE/POCKET NUMBER: 11746/1

TELEPHONE: (212) 425-7200

TELEPHONE: (212) 425-7200

TELERAX: (212) 425-5288

INPORMATION FOR SEQ ID NO: 16:
SEQUENCE CHRACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                               PRIOR APPLICATION DATA: APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 HWDFAWPW 8
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Gaps

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Gaps
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| Sequence 10, Application US/09794529
| Publication No. US20030082197A1
| GENERAL INFORMATION:
| APPLICANT: Sloan-Kettering Institute for Cancer Research
| ROTHMAN, James E. HARTL, F. Ulrich
| HOE, Mee H. HOGHTON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
| TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
| NUMBER OF SEQUENCES: 30 | CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 64; DB 3; Length 20; ilarity 100.0%; Pred. No. 0.071; Conservative 0; Mismatches 0: Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                   APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEPAX: (212) 425-7200
TELEPAX: (212) 425-7200
TELEPAX: (212) 425-7200
TELEPAX: (212) 425-7200
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FILING DATE: 27-Feb-2001
CLASSIFFCATION: Chassiffcation Date: Us/09/011,645
FILING DATE: 13-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20
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Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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US-09-794517-19
US-09-794517-19
Sequence 19, Application US/09794517
Publication No. US20030021794A1
GENERAL INFORMATION:
HAPLICANT: Sloan-Kettering Institute for Cancer Research
HARTL, F. Ulrich
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MARKEM, MARK
HUUGHTON: Heat Shock Protein-Based Vaccines and
Immunotherapies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION CURRENT: 27-Feb-2001
CLASSIFICATION: CURROWn>
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100.0%; Score 64; DB 3; Length 20;

Best Local Similarity 100.0%; Pred. No. 0.071;

Matches 8; Conservative 0; Mismatches 0; Indels
                   FILING DATE: <UNKNOWN:
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: AUGUST 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,439
REFERENCE/DOCKET NUMBER: 11/16/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                            TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: Delancia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECHANICALION INFORMATION:
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
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TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-528
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FILING DATE: 27-Feb-2001
GLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
sequence Description: Seq ID NO: 11: US-09-794-529-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGIAAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READMAILE FORM:
MEDIUM TYPE: Diskette - 3
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: MS DOS
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 11, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT:
ROTHMAN, James E.
HARTL, F. Ulrich
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0: Indels
                                                                              APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD POETECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIPICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                       NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELERAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 HWDFAWPW 20
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US-09-794-529-11
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
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                                                                                                                                                                          FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: Adgust 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
                                                                                                                                                   APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-794-529-13
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ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: yes FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                           PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT: GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION:
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: Diskette - 3.5 inch, 1.44 Mb storage
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COMPUTER: Diskette 
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APPLICATION NUMBER: US/c.,
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28, 839
REFERENCE/DOCKST NUMBER: 11746/1
TELEPHONE: (212) 425-5288
TELEPHONE: (212) 425-5288
TELEPAX: (212) 425-5288
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SEQUENCE DESCRIPTION: SEQ ID NO: 12:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: One Broadway
CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
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RESULT 139
US-09-744-529-16
Sequence 16, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
ROTHWAN, James E.
HARTL, F. Ulrich
HORE, Mee H.
HOUGHNON, Alan
TAKECHI, Yoshizumi
MATHEW, Mark
MATHEW, Mark
HOUGHNON, Mark
HOUGHNON, Mark
HOUGHNON, Mark
HOUGHNON, Mark
HOUGHNON, Mark
HOUGHNON, Mark
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                              PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
NAME: Defuncia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1000 A25-5288
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SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown>
                     CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
HARTL, F. Ulrich
HOB, Mee H.
HOUGHTON, Allan
TAKECHI, YOSHIZUMi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZID. 2 10004
COMPUTER F. 10004
COMPUTER F. 20004 FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                           FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                          NAME: Delucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEFAX: (212) 425-5288
TELEFAX: (210) 425-5288
TELEFAX: COLMINOWIN:
INFORMATION FOR SEQ 1D NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                  PRIOR APPLICATION DATA: APPLICATION NUMBER: US/09/011,645
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OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
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Best Local Similarity 100.
Matches 8; Conservative
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STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13 HWDFAWPW 20
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Gaps

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TYPE: amino acid
STRANDENESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: * < Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 30
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/09794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HARIL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, YOSHIZUMi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                      FRICAL MARCHES 13-F6b-1998
FILING DATE: 13-F6b-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
RAME: DELACIA, RICHARD I.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 28,839
TELECOMMUNICATION NUMBER: 11746/1
APPLICATION NUMBER: US/09/794,529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
               FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGNENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 30
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13 HWDFAWPW 20
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US-09-794-529-17
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich HOE, Mee H. HOUGHON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MUTUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 3; 100.0%; Pred. No. 0.071;
                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: Defucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus vaccine, seq ID NO: 17: US-09-794-529-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STRET: One Broadway
CITY: New York
STATE: NY
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-794-529-18
; Sequence 18, Application US/09794529
; Publication No. US20030082197A1
; GENERAL INFORMATION:
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RESULT 143
US-09-794-832-10
| US-09-794-832-10
| Sequence 10, Application US/09794832
| Publication No. US20030082198A1
| Publication No. US20030082198A1
| GENERAL INFORMATION:
| APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HORS, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi TAKECHI, Yoshizumi MAYMEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
100.0%; Score 64; DB 3;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0.
                     SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 61/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 18, 1995
APPLICATION NUMBER: 18, 1995
ATTORNEY/AGENT INFORMATION:
NAWE: DeLACIA REPERANCE, 1996
TELECOMMUNICATION NUMBER: 28, 839
REFERENCE/DOCKET NUMBER: 28, 839
REFERENCE/DOCKET NUMBER: 21, 1746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGWENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19, Application US/08794529
Publication No. US20030082197A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,529
FILING DATE: 27-Feb-2001
CLASSIFICATION: «Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
RILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
RILING DATE: August 16, 1996
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                               NAME: Defaucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEFAX: (212) 425-7200
TELEFAX: (212) 425-5288
TELEFAX: (212) 425-528
TELEFAX: COLORNOWID:
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunotherapies
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ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STRATE: NY
COUNTRY: US
ZIP: 10004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
    - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-F6-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
                       COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus vaccine...yutuu peptu ja seguence Description: Seguence Description: Seg ID NO: 11: US-09-794-832-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                              NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDENESS: «Unknown»
TOPOLGSY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12, Application US/09794832
Publication No. US20030082198A1
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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ZIP: 10004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11, Application US/09794832
Publication No. US20030082198A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOG, Mee H.
HOUGHTON, Alan
TAKECH! Yoshizumi
MATHEW, Mark Mark
MATHEW, Mark Mark
MATHEW, MARK MARK
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REGISTRATION NUMBER: 28,839
TELECOMMUNICATION INFORMATION:
COMPUTER: IBM compatible
OPERATING SYSTEMS, MS DOS
SOFTWARE: WORD Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-794-832-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunotherapies
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ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLGSY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10004
COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
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OTHER INFORMATION: hybrid peptide for human papilloma
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ROTHWAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
    ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                 COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF THE COMPUTER OF TH
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SEQUENCE DESCRIPTION: SEQ ID NO: 13:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-794-832-14
; Sequence 14, Application US/09794832
; Publication No. US20030082198A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: Yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: One Broadway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HWDFAWPW 8
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Publication No. US20030082198A1
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ultich
HOGHTON, Alan
TAKECHI, Yoshizumi
MATHEW, Mar B.
HOUGHTON, Alan
TAKECHI, Yoshizumi
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COMPUTER READABLE FORM:

MEDIUM TYPER Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFTCATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 64; DB 3; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  DATE APPLICATION DATA:
PAPLICATION NUMBER: US/09/011,645
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 18, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28, 939
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-5288
TELEPAX: (212) 425-5288
TELEPX: (212) 425-5288
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-09-794-832-12
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ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORIGINAL «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE
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Gaps

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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James B. HATL, F. Urich HOE, Mee H.
                                                                         - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <u >Umatemin Comparing Company</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-794-832-16; Sequence 16, Application US/09794832; Publication No. US20030082198A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
                      ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 30
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PEATURE
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US-09-794832-15
15-09-794832
1 Sequence 15, Application US/09794832
1 Publication No. US20030082198A1
1 GENERAL INFORMATION:
1 APPLICANT: Sloan-Kettering Institute for Cancer Research
1 HARTL, F. Ulrich
1 HARTL, F. Ulrich
2 HARTL, F. Ulrich
3 HARTEN, Noshizumi
4 MAYERM, Mark
5 TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
1 Immunotherapies
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                                                Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/1363
FILING DATE: AUGUST 16, 1996
ZIP: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.
COMPUTER: 1BM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
FILING DATE: 27-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <UNIONALION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS: LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18, Application US/09794832
Publication 0. US20030082198A1
GENERAL INFORMATION
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HOR, Mee H.
HOB, Mee H.
HOR, Mee H.
AUGGHTON, Alan
TARECHI, Yoshizumi
MAYHEW, Mark
                                                                                ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                     COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION NUMBER: US/09/011,645
FILING DATE: 13-Feb-1998
APPLICATION NUMBER: G0/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,499
FILING DATE: August 18, 1995
APPLICATION NUMBER: 61/995
APPLICATION NUMBER: B1,1995
APPLICATION NUMBER: B1,1995
APPLICATION NUMBER: B1,1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: DeLucia, Richard L.
REGISTRATION NUMBER: 28,839
REFRENCE/DOCKET NUMBER: 11746/1
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 30 CORRESPONDENCE ADDRESS:
                   CITY: New York
STATE: NY
                                                                  ns
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 HWDFAWPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HWDFAWPW 8
                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 151
US-09-794-832-18
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US-09-794-832-17

US-09-794-832-17

Sequence 17, Application US/09794832

Publication No. US20030082198A1

GENERAL INFORMATION:

ROTHWAN, Sames E. HARTL, F. Ulrich
HARTL, F. Ulrich
HOUGHOW, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark

MAYHEW, Mark

MAYHEW, Mark

MAYHEW, Mark

HOUGHOW, Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
NUMBER OF SEQUENCES: 30
                                         COUNTRY: US
CONTRY: US
COMPUTER: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
100.0%; Score 64; DB 3; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                            PRILICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION SATISTICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/011,645
FILING DATE: ABOUGT 19.8
APPLICATION NUMBER: 60/002,499
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 61/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: AUGUST 16, 1996
ATTORNEY/AGENT INFORMATION:
NAME: DELACIA, RICHARD L.
RECISTRATION NUMBER: 11,746/1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 425-5288
TELEX: «UNKOWN»
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 HWDFAWPW 20
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PRIOR APPLICATION TO COLALIDATE

APPLICATION NUMBER: US/09/011,645

PILING DATE: 13-F6-1998

APPLICATION NUMBER: 60/002,479

FILING DATE: August 18, 1995

APPLICATION NUMBER: 60/002,490

FILING DATE: August 18, 1995

APPLICATION NUMBER: PCT/US96/13363

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 28,839
REFERENCE/DOCKET NUMBER: 11746/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              yirus vaccine sequence bescription: SEQUENCE DESCRIPTION: SEQ ID NO: 19: US-09-794-832-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunotherapies
                        ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: DeLucia, Richard L.
                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM: WS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 425-5288
                                                                                                                          ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                            STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: yes
PRAGMENT TYPE: internal
ORIGINAL SOURCE:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                CITY: New York
                                                                                                        COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-170-713A-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19, Application US/09794832
Publication No. US20030082198A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Allan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                      ZIP: 10004
UTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                        PILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
                                                                                                                                                                  COMPUTER: IBM compatible
OPERATING SYSTEM; MS DOS
SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/794,832
FILING DATE: 27-Feb-2001
CLASSIPICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA: APPLICATION NUMBER: US/09/011,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 11746/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 18:
US-09-794-832-18
                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 13-Feb-1998
APPLICATION NUMBER: 60/002,479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 28,839
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: DeLucia, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDENESS: cubknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 425-5288
                    STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                  COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOGHTON, Alan
TAKEGH, YOSHİZUMİ
MAYHEGH, YOSHİZUMİ
MAYHEGH, YOSHİZUMİ
MAYHEGH, WARK
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
- 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 3; Length 20; 100.0%; Pred. No. 0.071;
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Gaps
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US-10-713A-12
S-G-10-713A-12
S-G-10-713A-12
S-G-10-10-713A-12
S-G-10-70-713A-12
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                                                                                                                                                                                                                                                                                                                                            - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.4
COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: IBM compatible
COMPUTER: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-7un-2002
CLASSIFICATION: 424
PRIOR APPLICATION ATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: August 18, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 00/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 19,1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               virus vaccine;
sequence description: seq ID NO: 11:
US-10-170-713A-11
       Immunotherapies
                     NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: Yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
ROTHMAN, James E.
HARTL, F. Ulrich
HOGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mash
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                  ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 64; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Caeson, Lawrence P.
REGISTATION NUMBER: 46,606
REFRENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: PCT/0896/1363
FILING DATE: AUGUST 16, 1996
                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JEQUENCE CHANCE.

JEQUENCE CHANCE.

LENGTH: 20

TYPE: amino acid
STRANDEDENESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGNETT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                               CITY: New York
STATE: NY
                                                                                                                                                                                                         COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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US-10-170-713A-11
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Gaps
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  MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
APPLICANT: SOJEMAN, James E.
HARTL, F. Ulrich
HOUGHTON, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.4

COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTMARE: Word Perfect
CURRENT APPLICATION DATA:
FILING DATE: 13-Jun-2002
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: February 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 09/011,645
                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                      STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
                                                                                                                                                              CITY: New York
                                                                                                                                                                                                           COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 157
US-10-170-713A-14
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: INDM COMPUTED DIskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/170,713A FILING DATE: 13-70n-2002 CLASSIFICATION DATA:

APPLICATION NUMBER: 09/011,645 FILING DATE: February 13, 1998 APPLICATION NUMBER: 60/002,479 FILING DATE: August 18, 1995 APPLICATION NUMBER: 60/002,490 FILING DATE: August 18, 1995 APPLICATION NUMBER: EVI/US96/13363 FILING DATE: August 18, 1995 APPLICATION NUMBER: PCT/US96/13363                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1:0

105-10-170-713A-13

1 Sequence 13, Application US/10170713A

2 Publication No. US20030185842A1

3 GENERAL INFORMATION:

4 APPLICANT: ROTHMAN, James E.

5 HARTL, F. Ulrich

6 HOUGHTON, Alan

7 TAKECHI, Yoshizumi
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 64; DB 4; Length 20; nlarity 100.0%; Pred. No. 0.071; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                         Immunotherapies
                                                               CORRESPONDENČE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFRAX: (212) 425-5288
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLGGY: linear
WOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                            NUMBER OF SEQUENCES: 30
                                                                                                                                                                                   COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 HWDFAWPW 20
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-170-713A-12
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HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
COUNTY: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
OTHER INFORMATION: hybrid peptide for human papilloma;
i virus vaccine,
;
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
US-10-713A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
COMPUTER: 10004
COMPUTER FORMS
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IDBACCLE 5.5.5 LICH, 1.3
COMPUTER: IDBACCOMPACTION
SOFTWARE: WORD PERFOCE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
CLASSIFICATION A24
PRIOR APPLICATION A24
PRIOR APPLICATION AVMBER: 09/011,645
FILING DATE: Rebruary 13, 1998
APPLICATION NUMBER: 60/002,499
FILING DATE: ANGUSE 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: 61/0502,490
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: 61/0502,490
FILING DATE: AUGUSE 18, 1995
APPLICATION NUMBER: 61/0502,490
FILING DATE: AUGUSE 16, 1995
APPLICATION NUMBER: PCT/USS6/13363
FILING DATE: AUGUSE 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: <UDACOMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: <Unknown>
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US-10-170-713A-16
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0
                                                                                                                                                    TAKECHI, Yoshizumi
MAXHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 150
US-10-170-713A-15
US-10-170-713A-15
Sequence 15, Application US/10170713A
Publication No. US20030185842A1
GENERAL INFORMATION:
HOPLICANT: Sloan-Kettering Institute for Cancer Research
HARTL, F. Ulrich
HARTL, F. Ulrich
HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 64; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REPERENCE DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAR: (212) 425-5288
TELEFAR: (1212) 425-5288
TELEFAR: (1212) 425-5288
TELEFAR: (1212) 425-7200
TELEFAR: (1212) 425-7200
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TELEFAR: (1212) 425-7200
TELEFAR: (1212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
CLASSIFICATION: 424
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SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13 HWDFAWPW 20
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Gaps

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Gaps
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                                                                                                           TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-170-713A-18

Sequence 18, Application US/10170713A

Publication No. US20030185842A1

GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                      - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h Similarity 100.0%; Score 64; DB 4; Length 20; Similarity 100.0%; Pred. No. 0.071; B; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                               COMPUTER: 1BM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: February 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
FILING DATE: August 18, 1995
FILING DATE: August 16, 1995
FILING DATE: August 16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
                                                                                                                                                                     CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
STREET: One Broadway
CITY: New York
STATE: NY
                                    HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                       NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                            ZIP: 10004
COMPUTER READABLE FORM:
                         HOE, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FELEX: <Unknown>
                                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 8; Conserva
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                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHMAN, James E.
                                                                                                                                                                                                                                                                               ZIP: 10004

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Casson, Lawrence P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/002,479
FILLING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
ORNEY/AGENT INPORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; virus vaccine
; SEQUENCE DESCRIPTION: SEQ ID NO: 16:
US-10-170-713A-16
                                                                                                              Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 09/011,645
                                                                                                                            NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17, Application US/10170713A Publication No. US20030185842A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
                    HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRACHENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                   COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13 HWDFAWPW 20
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MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
Sloan-Kettering Institute for Cancer Research ROTHMAN, James B. HARTL, F. Ulrich HOE, Mee H. HOUGHTON, Alan TAKECHI, Yoshizumi
                                                                                                                                                                                                                                                                                                                                                                           - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: INFECTOR OF STATES, CARE OF STEM, MS DOS SOFTWARE WAS DOS SOFTWARE WAS DELECT CURRENT APPLICATION DATA:

APPLICATION DATA:
APPLICATION NUMBER: US/10/170,713A
FILING DATE: 13-Jun-2002
CLASSIFICATION NUMBER: 09/011,645
APPLICATION NUMBER: 09/011,645
APPLICATION NUMBER: 09/012,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 00/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
ATORNEY/AGENT INFORMATION:
NAME: CASSON, LAWFENCE P.
REGISTRATION NUMBER: 46,606
RESTERATION NUMBER: 46,606
RESTERATION NUMBER: 46,606
RESTERATION NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
TELESPHONE: (212,425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               y virus vaccine ;
seQUENCE DESCRIPTION: SEQ ID NO: 19: US-10-170-713A-19
                                                                                                                                                                                 NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
PRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                              ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                       CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: hybrid peptide for human papilloma
    ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark MAHEW, Mark Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: Liskette - 3.5 inch, 1.44 Mb storage Deraylus System: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:

FILING DATE: 13-Jun-2002
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Casson, Lawrence P.
REGISTATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 18, 1995
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virus vaccine
sequence Description; seq ID NO: 18:
US-10-170-713A-18
                                                                                                                                                   Immunotherapies
                                                                                                                                                                                                    ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 162
US-10-170-713A-19
; Sequence 19, Application US/10170713A
; Publication No. US20030185842A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 425-72(
TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                    NUMBER OF SEQUENCES: 3(CORRESPONDENCE ADDRESS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                              COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 HWDFAWPW 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ઠે
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OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                            100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                        US-10-171-734-10; Sequence 10, Application US/10171734; Publication No. US20030185843A1
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OTHER INFORMATION: hybrid peptide for human papilloma
                   Sequence 11, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKEGHI, Yoshizumi
MAYHEW, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IDBRECTEE - 3.5 inch, 1.44 |
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: August 10, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
ATTORNEY/AGENT INFORMATION:
NAME: CASSON, LAWFENCE P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION NUMBER: 11746/461031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSES Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE
  US-10-171-734-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-171-734-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                    TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                     Sloan-Kettering Institute for Cancer Research ROTHMAN, James E. HARTL, F. Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                 - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Casson, Lawrence P.
REGISTATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/1363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: Pebruary 13, 1998
APPLICATION NUMBER: 09/616,295
FILING DATE: August 10, 2000
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTHARR: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/171,734
FILLING DATE: 14-Jan-2003
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                     HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: «Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: «Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (212) 425-5288
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                                                                                                                                                                                                                                                                                                 STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FELEPHONE: (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HWDFAWPW 20
GENERAL INFORMATION
                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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Gaps

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RESULT 164

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Sequence 13, Application US/10171734

Publication No. US20030185843A1

GENERAL INFORMATION:

APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHAND, James E.

HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immnotherapies

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESSE:
ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway

CITTY: New York

CITTY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; DB 4; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: INSECUE.

COMPUTER: INSECUE.

SOFTWARE: WACD PERFECT

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/171,734

FILING DATE: 14-Jan-2003

CLASSIFICATION NUMBER: 60/002,479

FILING DATE: August 18, 1995

APPLICATION NUMBER: 60/002,490

FILING DATE: August 18, 1995

APPLICATION NUMBER: 00/01,645

FILING DATE: August 16, 1996

APPLICATION NUMBER: 09/011,645

FILING DATE: August 16, 1996

APPLICATION NUMBER: 09/011,645

FILING DATE: August 10, 2000

APPLICATION NUMBER: 09/636,295

FILING DATE: August 10, 2000

ATTORNEY AGENT INFORMATION:

NAME: CASSON, LAWICHORY

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yea
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
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13 HWDFAWPW 20
HWDFAWPW 8
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                                                                                                                      RESULT 166
US-10-171-734-13
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MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION DATA:
APPLICATION NUMBER: 06/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 09/011,645
FILING DATE: Rebruary 13, 1998
APPLICATION NUMBER: 09/011,645
FILING DATE: Rebruary 13, 1998
APPLICATION NUMBER: 09/011,645
FILING DATE: August 10, 2000
ATTORNEY/AGENT INFORMATION:
NAME: CREBON LAWRENCE P.
REFERENCE/DOCKET NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                          Sequence 12, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James B.
HARIL, F. Ulrich
HOE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
RADBRESSER: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
TELEX: «UNKNOWN»
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: NY
                                                                           .0 - 171 - 734 - 12
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Gaps
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US-10-111-734-15
i Sequence 15, Application US/10171734
j Publication No. US20030185843A1
j GENERAL INFORMATION:
i GENERAL INFORMATION:
i HAPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James E.
HARTL, F. Ulrich
HOB, Mee H.
HORDHON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
MAYHEW, Mark
Heat Shock Protein-Based Vaccines and
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                                               100.0%; Score 64; DB 4; Length 20; ilarity 100.0%; Pred. No. 0.071; Conservative 0; Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: 90/011,645
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDFLUAY 13, 1998
APPLICATION NUMBER: 09/011,645
FILING DATE: August 10, 2000
ATTORNEY/AGENT INPORMATION:
NAME: CASSON, LAWFENCE P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM compatible
OPERATING SYSTEM
SOFTWARE: Word Parfect
CURRENT APPLICATION DATA:
RELING DATE: 14-Jan-2003
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 425-5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30 CORRESPONDENCE ADDRESS:
                                                                                                                                                                             13 HWDFAWPW 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                              1 HWDFAWPW 8
                                                        Query Match
Best Local Similarity
Matches 8; Conserv
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             US-10-171-734-14
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                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
                                                                                                                                                                                                                             Sequence 14, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHMAN, James B.
HARTL, F. Ulrich
HOGE, Mee H.
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAXHEW, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: MS DOS SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDRUARY 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
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REFERENCE/DOCKET NUMBER: 11746/461031
TELECOMMUNICATION INFORMATION:
                  Pred. No. 0.(
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SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Casson, Lawrence P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
100.08; F1.
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STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: <Unknown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10004
COMPUTER READABLE FORM:
               Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCES:
                                                                                 HWDFAWPW 8
                                                                                                              HWDFAWPW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE
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TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
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  ORIGINAL SOURCE:
ORGANISM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                 13 HWDFAWPW 20
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                                          PEATURE:
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                                                                                                                                                       Gaps
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18-10-71-734-16

1 Sequence 16, Application US/10171734

2 Publication No. US20030185843A1

3 GENERAL INFORMATION:

4 APPLICANT: Sloan-Kettering Institute for Cancer Research

5 HARTL, F. Ulrich

6 HARTL, F. Ulrich

7 HOUG, Mee B.

1 HOUGHTON, Alan

7 TAKECHI, Yoshizumi

MAYHEW, Mark

MAYHEW, Mark

1 AAKECHI, Yoshizumi

MAYHEW, Mark

1 AAKECHI, Yoshizumi

1 AAKECHI, Shock Protein-Based Vaccines and
                                                                                                                                                       ò
OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                        Score 64; DB 4; Length 20; Pred. No. 0.071;
                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/002,479
FILING DATE: ANGUER 18, 1995
APPLICATION NUMBER: 60/002,490
FILING DATE: ANGUER 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 09/011,645
FILING DATE: PEDTUARY 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: ANGUER 10, 2000
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: CASBON, LAWYERCE P.
REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
COMPUTER: 1BM compatible
COMPATING SYSTEM: MS DOS
SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
RAPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIPICATION: 424
                                                                                                                                                   0; Mismatches
                      virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSE: Kenyon & Kenyon STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212) 425-7200
TELEFAX: (212) 425-5288
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TELEPHONE: (212) 425-720(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: «Unknown»
TOPOLOGY: linear
MOLECULE TYEE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
                                                                                                        Query Match
Best Local Similarity 100.0%;
Matches 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
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1 HWDFAWPW 8
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                                                              US-10-171-734-15
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13 HWDFAWPW 20
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                                                                                                                           OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sloan-Kettering Institute for Cancer Research ROTHWAN, James E. HARTL, F. Ulrich HOE, Mee H. HOEHON, Alan TAKECHI, Yoshizumi MAYHEW, Mark
                                                                                                                                                                                                                                   Query Match 100.0%; Score 64; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                             Indels
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APPLICATION NUMBER: PCT/US96/13363
FILING DATE: August 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10004

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1
COMPUTER: IBM compatible
OPERATING SYSTEM: MS DOS
SOFTWARE: WORD PETFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 17:
US-10-171-734-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/002,479
FILING DATE: AUGUST 18, 1995
APPLICATION NUMBER: 60/002,490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunotherapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSES: Kenyon & Kenyon
STREET: One Broadway
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Casson, Lawrence P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18, Application US/10171734
Publication No. US20030185843A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
                                                                                      ORGANISM: <Unknown>
MOLECULE TYPE: peptide
HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA
                                                                                                                                                   virus vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                       1 HWDFAWPW 8
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                                                                                                   FEATURE
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US-10-17-734-19
; Sequence 19, Application US/10171734
; Publication No. US20030185843A1
; GENERAL INFORMATION:
; APPLICANT: Sloan-Kettering Institute for Cancer Research
ROTHWAN, James E.
HARTL, F. Ulrich
HARTL, F. Ulrich
HOUGHTON, Alan
TAKECHI, Yoshizumi
MAYHEW, Mark
; TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and
Immunotherapies
                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                       OTHER INFORMATION: hybrid peptide for human papilloma
                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 64; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 46,606
REFERENCE/DOCKET NUMBER: 11746/461031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/002,479
FILING DATE: August 18, 1995
PPLICATION NUMBER: 60/002,490
FILING DATE: August 18, 1995
APPLICATION NUMBER: PCT/US96/13363
APPLICATION NUMBER: 16, 1996
APPLICATION NUMBER: 09/011,645
FILING DATE: February 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/10/171,734
FILING DATE: 14-Jan-2003
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                  virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 09/636,295
FILING DATE: August 10, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 425-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Casson, Lawrence P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM compatible
OPRRATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: One Broadway
CITY: New York
STATE: NY
                                                                                                                                          ORGANISM: <Unknown>
                                                                  HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
ZIP: 10004
COMPUTER READABLE FORM:
MEDIUM TYPE: Diske
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Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: javelinized peptide of gp100 human melanoma OTHER INFORMATION: antigen
                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071;
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Sequence 26, Application US/10258147
Publication No. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Rothman, James
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REPRENCE: 11746/4627
CURRENT APPLICATION UNBER: US/10/258,147
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rothman, James
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,462
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEG ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 25
LENGTH: 20
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                              virus vaccine
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
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PRIOR APPLICATION NUMBER: 60/197,462
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                                                                                                                                                   HYPOTHETICAL: yes
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
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Matches 8; Conservative
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1 HWDFAWPW 8
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US-10-258-147-25
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GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich

APPLICANT: Hodyhoon, Alan

APPLICANT: Hodyhoon, Alan

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Maybew, Mark

TITLE OF INVENTION: Hat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461061

CURRENT APPLICATION NUMBER: US/10/367,580

CURRENT FILING DATE: 2003-02-14

PRIOR APPLICATION NUMBER: US 09/794,832

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR APPLICATION NUMBER: US 60/002,479

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

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                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: javelinized peptide of human gp100 melanoma; OTHER INFORMATION: antigen
US-10-258-147-26
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 26
LENGTH: 20
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US-10-367-580-10
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APPLICANT: Rothman, James E.; APPLICANT: Hartl, F. Ulrich; APPLICANT: Hoe, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-367-580-10
; Sequence 10, Application US/10367580
; Publication No. US20040071720A1
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ORGANISM: Artificial Sequence
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APPLICANT: Rothman, James B.
APPLICANT: Harll, F. Ulrich
APPLICANT: Harll, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Houghton, Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
CURRENT APPLICATION NUMBER: US/10/367,580
CURRENT PILING DATE: 2003-02-14
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APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
IIILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
           Indels
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           Mismatches
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PRIOR FILING DATE: 2001-02-27
PRIOR PELLING DATE: 2001-02-27
PRIOR PELLING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 13
LENGTH: 20
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CURRENT APPLICATION NUMBER: US/10/367,580
CURRENT FILING DATE: 2003-02-14
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APPLICATION NUMBER: US 09/011,645
FILING DATE: 1998-02-13
APPLICATION NUMBER: PCT/US96/13363
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PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
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; OTHER INFORMATION: synthetic peptide
US-10-367-580-13
                                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/10367580 Publication No. US20040071720A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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           8; Conservative
                                                                                                                               13 HWDFAWPW 20
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APPLICANT: Hartl, F. Ulrich
APPLICANT: How, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE ON INVENTION: Has Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461061
                                                                                                         TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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Pred. No. 0.071;
                                                                                                                                                                                                          CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,832
PRIOR FILING DATE: 2003-02-14
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR PILING DATE: 1996-08-16
PRIOR PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOOTWARE: WordPerfect 8.0 for Windows SEQ ID NO 11
LENGTH: 20
TYPE: PRI

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PRIOR FILING DATE: 2001-02-27
PRIOR PELING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1998-02-13
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTUM REI WORDPERE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SSOFUMARE: WordPerfect 8.0 for Windows
LENGTH: 20
                                                                                                                               FILE REFERENCE: 11746/461061
CURRENT APPLICATION NUMBER: US/10/367,580
CURRENT FILING DATE: 2003-02-14
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; OTHER INFORMATION: synthetic peptide
US-10-367-580-12
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Houghton, Alan
Takechi, Yoshizumi
Mayhew, Mark
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 9; Conservative
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Best Local Similarity
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1 HWDFAWPW 8
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LENGTH: 20
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Sequence 15, Application US/10367580

Publication No. US20040071720A1

GENERAL INPORMATION:

APPLICANT: Hothman, James E.

APPLICANT: Hothcon, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Mark

TILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461061

CURRENT PRILING DATE: 2003-02-14

PRIOR APPLICATION NUMBER: US 09/194,832

PRIOR APPLICATION NUMBER: US 09/11,645

PRIOR PLING DATE: 1998-02-13

PRIOR PLING DATE: 1998-02-13

PRIOR PLING DATE: 1995-08-18

PRIOR PLING DATE: 1995-08-18

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PRIOR FILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SOFTWARE: WordPerfect 9.0 for Windows
LENGTH: 20
TYPE: PRT
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; OTHER INFORMATION: synthetic peptide
US-10-367-580-15
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Sequence 16, Application US/10367580
Publication No. US20040071720A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461061
CURRENT PILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,832
PRIOR APPLICATION NUMBER: US 09/711,645
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR PILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
NUMBER OF SEQ ID NOS: 349
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APPLICANT: Harly, F. Ulrich
APPLICANT: Hote, wee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Advhew, Mark
ITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/10/367,580
CURRENT FILING DATE: 2003-02-14
PRIOR PELLING DATE: 2003-02-17
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SEQ ID NO 17
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: synthetic peptide
US-10-367-580-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
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Gaps

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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461012
CURRENT APPLICATION NUMBER: US/10/367,593
CURRENT FILING DATE: 1998-02-14
PRIOR FILING DATE: 1998-02-13
PRIOR PELING DATE: 1998-02-13
PRIOR PELING DATE: 1995-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
ESEQ ID NO 19
LENGTH: 20
                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: synthetic peptide US-10-367-580-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: synthetic peptide US-10-367-593-10
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Publication No. US20040071721A1
GENERAL INFORMATION:
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Publication No. US20040071721A1
GENERAL INFORMATION:
                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich APPLICANT: Hoe, Mee H.
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Publication No. US20040071720A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Rothman, James B.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Mayhew, Mark

TILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

CURRENT APPLICATION NUMBER: US/10/367,580

CURRENT FILING DATE: 2003-02-14

FRIOR PELING DATE: 2001-02-27

PRIOR APPLICATION NUMBER: US 69/011,645.

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,479

PRIOR APPLICATION NUMBER: US 60/002,479

PRIOR APPLICATION NUMBER: US 60/002,479

PRIOR FILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

SOFTWARE: WordPerfect 8.0 for Windows

LENGTH: 20

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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461061
CURRENT APPLICATION NUMBER: US/10/367,580
                                  Gaps
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                                  Indels
     Pred. No. 0.071;
                                  Mismatches
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PRIOR FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR FILING DATE: 1998-02-13
PRIOR PLICATION NUMBER: PCT/US96/13363
PRIOR FILING DATE: 1996-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: synthetic peptide
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        100.08;
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ORGANISM: Artificial Sequence
     Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
Matches 8; Conserv
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TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 100. ...
8; Conservative
        13 HWDFAWPW 20
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Sequence 12, Application US/10367593

Publication No. US20040071721A1

GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Alan

APPLICANT: Takechi, Yoshizumi

APPLICANT: Houghton, Alan

APPLICANT: Takechi, Yoshizumi

APPLICANT: Town Warker, US/10/367,593

CURRENT APPLICATION NUMBER: US/10/367,593

CURRENT FILING DATE: 1998-02-13

PRIOR APPLICATION NUMBER: US/10/36/13363

PRIOR APPLICATION NUMBER: US/09/01,645

PRIOR PILING DATE: 1995-08-16

PRIOR FILING DATE: 1995-08-16

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

SOFTWARE: WordPerfect B.O for Windows

SEQ ID NO 12

LENGTH: 20

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SEQ ID NO 12

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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mark
TITLE OF INVENTION: Mark
TITLE OF INVENTION: Mark
FILE REFERENCE: 11746/461012
CURRENT APPLICATION NUMBER: US/10/367, 593
CURRENT FILING DATE: 1989-02-14
PRIOR APPLICATION NUMBER: US/0011,645
PRIOR APPLICATION NUMBER: US/0011,645
PRIOR APPLICATION NUMBER: US/002,490
PRIOR PILING DATE: 1995-08-16
PRIOR APPLICATION NUMBER: US/0002,490
PRIOR PILING DATE: 1995-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 11
LENGTH: 20
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) OTHER INFORMATION: Synthetic peptide US-10-367-593-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich APPLICANT: Hoe, Mee H.
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APPLICANT: Hartl, F. Ulrich
APPLICANT: How, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4461012
CURRENT APPLICATION NUMBER: US/10/367,593
CURRENT FILING DATE: 2003-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16. Application US/10367593
Sequence 16. Application US/10367593
Septicant No. US20040071721A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yeshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461012
CURRENT APPLICATION VUMBER: US/10/367,593
CURRENT FILING DATE: 2003-02-14
                                                                                                                                                                          Gaps
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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PRIOR PELING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: WG 9/011,645
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER: OF SEQ ID NOS: 349
SEQ ID NO 15
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ON NUMBER: US 09/011,645
                                               ; OTHER INFORMATION: synthetic peptide US-10-367-593-14
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Publication No. US20040071721A1
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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HWDFAWPW 8
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PRIOR PAPLICATION NUMBER: PCT/USPA/13302
PRIOR PAPLICATION NUMBER: PCT/USPA/13302
PRIOR PAPLICATION NUMBER: 1956/002,490
PRIOR PAPLICATION NUMBER: 1956/002,490
PRIOR PAPLICATION NUMBER: 1956/002,479
PRIOR PAPLICATION NUMBER: 1956/002,479
PRIOR PAPLICATION NUMBER: 1956/002,479
PRIOR PAPLICATION NUMBER: 1956/002,479
PRIOR PAPLICATION NUMBER: 1956/002,479
PRIOR PAPLICATION NUMBER: 1966/002,479
PRIOR PAPLICATION SPACE AND SPACE 64, DB 4; Length 20;
PRESULT 192
PRIOR PAPLICATION SPACE 64, DB 4; Length 20;
PRESULT 192
PRIOR PAPLICATION SPACE 64, DB 4; Length 20;
PRESULT 192
PRIOR PAPLICATION NUMBER: 105/001/157/593
PRIOR PAPLICATION NUMBER: 105/001/597/593
PRIOR PAPLICATION NUMBER: 105/001/597/593
PRIOR PLICATION NUMBER: 1
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13 HWDFAWPW 20
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| GENERAL INPORMATION:
| APPLICANT: Hartl, F. Ulrich
| APPLICANT: Hore, Mee H.
| APPLICANT: Howehen, James E.
| APPLICANT: Howehen, James H.
| APPLICANT: Howehen, Alan
| APPLICANT: Takechi, Yoshizumi
| APPLICANT: Takechi, Yoshizumi
| APPLICANT: Takechi, Yoshizumi
| APPLICANT: Mark
| TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
| TILE REFERENCE: 11746/461012
| CURRENT APPLICATION NUMBER: US/10/367,593
| CURRENT PILING DATE: 1998-02-13
| PRIOR APPLICATION NUMBER: US 60/001,490
| PRIOR FILING DATE: 1995-08-16
| PRIOR FILING DATE: 1995-08-18
| PRIOR FILING DATE: 1995-08-18
| PRIOR FILING DATE: 1995-08-18
| NUMBER OF SEQ ID NOS: 349
| SEQ ID NO 19
| LENGTH: 20
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| LENGTH: 20
| LENGTH: 20
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE REFERENCE: 11746/461012
CURRENT APPLICATION NUMBER: US/10/367,593
CURRENT FILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-18
PRIOR PILING DATE: 1996-08-18
PRIOR PILING DATE: 1996-08-18
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SOFTWARE: WordPerfect 8.0 for Windows
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: synthetic peptide US-10-367-593-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , OTHER INFORMATION: synthetic peptide
US-10-367-593-19
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ORGANISM: Artificial Sequence
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TITLE OF INVENTY OF A PACEDIAN AND A PLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REPERBUES: 1176/461014
CURRENT APPLICATION NUMBER: US/10/367,594
CURRENT PILING DATE: 2003-02-14
FRIOR PELING DATE: 2000-10-05
FRIOR PAPLICATION NUMBER: US 09/011,645
FRIOR PILING DATE: 1998-02-13
FRIOR FILING DATE: 1998-02-13
FRIOR PILING DATE: 1998-02-13
FRIOR PILING DATE: 1995-08-16
FRIOR APPLICATION NUMBER: US 60/002,490
FRIOR PILING DATE: 1995-08-18
FRIOR APPLICATION NUMBER: US 60/002,479
FRIOR PILING DATE: 1995-08-18
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FRIOR PILING DATE: 1995-08-18
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FRIOR APPLICATION NUMBER: US 60/002,479
FRIOR FILING DATE: 1995-08-18
FRIOR APPLICATION NUMBER: US 60/002,479
FRIOR FILING DATE: 1995-08-18
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SOFTWARE: WORDPERFECT 8:0 for Windows
SEQ ID NO 13
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Publication No. US20040071722A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Hoc, Mee H:
APPLICANT: Hoc, Mee H:
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew; Mark
APPLICANT: Mayhew; Mark
APPLICANT: INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461041
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CURRENT APPLICATION NUMBER: US/10/367,594
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/680,806
PRIOR FILING DATE: 2000-10-05
PRIOR PILING DATE: 1996-02-13
PRIOR PILING DATE: 1996-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
....nhes 8; Conservative
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Publication No. US20040071722A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4610418.
CURRENT APPLICATION NUMBER: US/10/367,594
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PRIOR APPLICATION WUMBER: US 09/680,806
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WORDPEFFECE 8.0 for Windows
LENGTH: 20
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 11
LENGTH: 20
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                                                                                                                                                                                                                                                                                      , OTHER INFORMATION: synthetic peptide US-10-367-594-11
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Best Local Similarity 100.0%;
Matches 8; Conservative (
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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| Sequence 16, Application US/10367594
| Publication No. US20040071722A1
| GENERAL INFORMATION:
| APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich
| APPLICANT: Hartl, F. Ulrich
| APPLICANT: Houghton, Alan
| APPLICANT: Takechi, Yoshizumi
| APPLICANT: Mayhew, Mark
| TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
| FILE REFERENCE: 11746/461041
| CURRENT ELLING DATE: 2009-02-14
| PRIOR PILLING DATE: 2009-02-14
| PRIOR PELLING DATE: 1998-02-13
| PRIOR PILLING DATE: 1998-02-13
| PRIOR FILLING DATE: 1996-08-16
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APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Takechi, Yoshizum
APPLICANT: Takechi, Yoshizum
APPLICANT: Mayhew, Mark
ITITE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461041
CURRENT APPLICATION NUMBER: US/10/367,594
CURRENT FILING DATE: 2000-10-05
FRIOR PILING DATE: 1996-08-16
FRIOR PRILING DATE: 1996-08-16
FRIOR PILING DATE: 1996-08-16
FRIOR APPLICATION NUMBER: US 60/002,490
FRIOR FILING DATE: 1995-08-18
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; Sequence 15, Application US/10367594
; Publication No. US20040071722A1
; GENERAL INFORMATION:
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1 HWDPAWPW 8
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US-10-367-594-17

US-10-367-594-17

Sequence 17, Application US/10367594

Publication No. US20040071722A1

GENERAL INFORMATION:

APPLICANT: Horhman, James E.

APPLICANT: How, Mee H.

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Maybaw, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 1746/461041

CURRENT PILING DATE: 2000-10-14

CURRENT PILING DATE: 2000-10-05

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

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NUMBER OF SEQ ID NOS: 349

SOFTWARE: WORTCHECK 8.0 for Windows
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PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 16
LENGTH: 20
                                                                                                                                                                                                          ; FEATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-594-16
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APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461041
CURRENT APPLICATION NUMBER: US /10/367,594
CURRENT FILING DATE: 2003-02-14
PRIOR PRILING DATE: 2000-10-05
PRIOR PELING DATE: 2000-10-05
PRIOR PELING DATE: 1996-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR PILING DATE: 1995-08-18
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APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Howe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
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APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mark
TILE OF INVENTION: Haat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461041
CURRENT APPLICATION NUMBER: US/10/367,594
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/680,806
PRIOR PRILING DATE: 1998-02-13
PRIOR PRILING DATE: 1996-08-18
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; OTHER INFORMATION: synthetic peptide
US-10-367-594-19
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Publication No. US20040071722A1
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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APPLICANT: Rothman, James E.
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APPLICANT: Houghton, Alan
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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Maark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT FILING DATE: 2003-02-14
FRIOR APPLICATION NUMBER: US 10/171, 734
FRIOR PRICATION NUMBER: US 09/636, 295
FRIOR FILING DATE: 2000-08-10
FRIOR FILING DATE: 2000-08-10
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FRIOR APPLICATION NUMBER: US 09/011,645
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APPLICANT: Harl, F. Ulrich
APPLICANT: Hoe, Mee
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhww, Mark
APPLICANT: Mayhww, Mark
TITLE OF INVENTION: Hear Shock
FILE REFERENCE: 11746/461032
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PRIOR PILING DATE: 2002-06-13
PRIOR PILING DATE: 2002-06-13
PRIOR APPLICATION UNDRER: US 09/636,295
PRIOR FILING DATE: 2000-08-10
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-16
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PRIOR FILING DATE: 1995-08-18
SOFTHARE: WORDERFER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
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; OTHER INFORMATION: synthetic peptide
US-10-367-654-10
                                                                                                                                                                           Sequence 10, Application US/10367654 Publication No. US20040071723A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11, Application US/10367654
Publication No. US20040071723A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Matches 8; Conservative
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GABLEANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US 10/171,734
PRIOR APPLICATION NUMBER: US 10/171,734
PRIOR PILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,490
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR PLING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SSQTWARE: WordPerfect 8.0 for Windows
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Synthetic peptide US-10-367-654-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Synthetic peptide US-10-367-654-12
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Publication No. US20040071723A1
GENERAL INFORMATION:
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Mark
TITIE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US/10/367,654
CURRENT APPLICATION NUMBER: US 10/171,734
PRIOR FILING DATE: 2003-02-13
PRIOR PELING DATE: 2000-08-10
PRIOR PILING DATE: 1098-02-13
PRIOR PELING DATE: 1998-02-13
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PRIOR PELING DATE: 1995-08-18
NUMBER OF ERQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 17
LENGTH. 20
      APPLICANT: Targetat, Towareaut, APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US/10/367,654
CURRENT PILING DATE: 2003-02-14
PRIOR PILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 09/636,295
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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COTHER INFORMATION: synthetic peptide
US-10-367-654-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) OTHER INFORMATION: synthetic peptide US-10-367-654-16
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Publication No. US20040071723A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
   Takechi, Yoshizumi
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ORGANISM: Artificial Sequence
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APPLICANT: Rothman, James B.
APPLICANT: Hartl, F. Ulrich
APPLICANT: How, Mee H.
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US/10/367,654
CURRENT FILING DATE: 2003-02-14
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PRIOR FILING DATE: 2002-06-13
PRIOR PELING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 09/636,295
PRIOR FILING DATE: 2000-08-10
PRIOR PELING DATE: 1998-02-13
PRIOR PELING DATE: 1998-02-13
PRIOR PELING DATE: 1998-08-16
PRIOR PELING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
PRIOR PELING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
LENGTH: 20
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Publication No. US20040071723A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide
US-10-367-654-14
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SOFTWARE: WordPerfect 8.0 for Windows
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Publication No. US20040071723A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity الاست
الاستخداد 8، Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-367-658-10
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Sequence 18, Application US/10367654

Sequence 18, Application US/10361654

Sequence 18, Application US/10361654

SEDELICANT: Rethman, James E.

APPLICANT: Hart, F. Ulrich

APPLICANT: How, Mee H.

APPLICANT: Howelven, Alan

APPLICANT: Harthe, Mee H.

APPLICANT: Harthe, Mee H.

APPLICANT: Harthe, Mee H.

APPLICANT: Harthe, Mee H.

APPLICANT: Harthe, Mee H.

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/46103

CURRENT APPLICATION NUMBER: US/10/367,654

CURRENT PILING DATE: 2002-06-13

PRIOR FILING DATE: 1098-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-08-18

PRIOR FILING DATE: 1995-08-18

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APPLICANT: Harll, F. Ulrich
APPLICANT: Horline, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461032
CURRENT APPLICATION NUMBER: US/10/367,654
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 10/171,734
PRIOR PLING DATE: 2002-06-13
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  100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071; ive 0; Mismatches 0; Indels
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; OTHER INFORMATION: synthetic peptide
US-10-367-654-18
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Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                      1 HWDFAWPW 8
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1 HWDFAWPW 8
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Sequence 10, Application US/10367658

Sequence 10, Application US/10367658

Bublication No. US20040071724A1

GRNEAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
CURRENT APPLICATION NUMBER: US/10/367,658
CURRENT FILING DATE: 2003-02-14
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1995-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: US 09/636,295
PRIOR FILING DATE: 2000-08-10
PRIOR PPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTHARE: WORDPEFFECE 8:0 for Windows
SEQ ID NO 19
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                     ; FEATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-654-19
                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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                                                          SCHERAL INFUGATION:

APPLICANT: Rothman, James E.
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hack, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
ITILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
CURRENT APPLICATION NUMBER: US/10/367,658
CURRENT APPLICATION NUMBER: US 09/794,529
PRIOR FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1996-08-15
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SQCFTWARE: WordPerfect 8.0 for Windows
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTYON: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
CURRENT APPLICATION NUMBER: US/10/367,658
CURRENT FILING DATE: 2003-02-14
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PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 1998-02-13
PRIOR PELLING DATE: 1998-02-13
PRIOR PELLING DATE: 1996-09-16
PRIOR PELLING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
LENGTH: 20
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Sequence 11, Application US/10367658
Publication No. US20040071724A1
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Publication No. US20040071724A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
CURRENT APPLICATION NUMBER: US 09/794,529
FRIOR APPLICATION NUMBER: US 09/794,529
FRIOR APPLICATION NUMBER: US 09/011,645
FRIOR PELING DATE: 1996-08-16
FRIOR FILING DATE: 1995-08-16
FRIOR PELING DATE: 1995-08-18
FRIOR APPLICATION NUMBER: US 60/002,479
FRIOR APPLICATION NUMBER: US 60/002,479
FRIOR APPLICATION NUMBER: US 60/002,479
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APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mar
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/10/367,658
ORGANISM: Artificial Sequence
PEATURE:
JOTHER INFORMATION: Synthetic peptide
US-10-367-658-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 349
SOFWARE: WordPerfect 8.0 for Windows
SEQ ID NO 13
LENGTH: 20
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; OTHER INFORMATION: synthetic peptide
US-10-367-658-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13, Application US/10367658
Publication No. US20040071724A1
GENERAL INFORMATION:
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Publication No. US20040071724A1
GENERAL INFORMATION:
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
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US-10-367-658-17
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### APPLICANT: No. US20040071724A1

GENERAL INFORMATION:

### APPLICANT: Rothman, James E.

### APPLICANT: Harl, F. Ulrich

### APPLICANT: Hoe, Mee H.

### APPLICANT: Hoe, Mee H.

### APPLICANT: Mayhew, Mark

### TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

### TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

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### TILLE OF INVENTION: Heat Shock Protein-Based Vaccines

### TILLE OF INVENTION: Heat Shock Protein-Based Vaccines

### TILLE OF INVENTION: Heat Shock Protein-Based Vaccines

### CURRENT FILING DATE: 1908-02-13

### PRIOR FILING DATE: 1996-08-16

### PRIOR FILING DATE: 1996-08-16

### PRIOR FILING DATE: 1995-08-18

### PRIOR FILING DATE: 1995-08-18

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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION WUMBER: US 09/794,529
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION WUMBER: WG 00/002,490
PRIOR PILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WORDPERFECT 8.0 for Windows
LENGTH: 20
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; OTHER INFORMATION: synthetic peptide
US-10-367-658-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: synthetic peptide
US-10-367-658-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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us-10-20,-bus-10,
publication No. US2004007125A1
generce 10, Application US/10367668
gbublication No. US2004007125A1
generce 10, Application US/10367668
general invokation:
garpicant: Rothman, James E.
garpicant: Hoc, Mee H.
garpicant: Hoc, Mee H.
garpicant: Hoc, Mee H.
garpicant: Hoc, Mee H.
garpicant: Mayhew, Mark
garpicant: Mayhew, Mark
garpicant: Mayhew, Mark
garpicant: Mayhew, Mark
garpicant: Mayhew, Mark
garpicant: Mayhew, Mark
garpicant: 11746/461072
gurrant Pring Application Number: US/10/367,668
gurrant Applicanton Number: US 09/794,517
grior Riling Date: 1098-02-13
grior Filing Date: 1996-08-16
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CURRENT APPLICATION NUMBER: US/10/367,658
CURRENT FILING DATE: 2003-02-14
PRIOR PELLING DATE: 2003-02-37
PRIOR FILING DATE: 2001-02-27
PRIOR PLING DATE: 1998-02-13
PRIOR PLING DATE: 1998-02-13
PRIOR PLING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: PCT/USS6/13363
PRIOR FILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SEQ THARB: WARDPERFECT B.O. FOR WINDOWS
SEQ THARBER: WARDPERFECT B.O. FOR WINDOWS
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; OTHER INFORMATION: synthetic peptide
US-10-367-658-19
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; OTHER INFORMATION: synthetic peptide
US-10-367-668-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 8; Conservative
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APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhaw, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 1746/461051
CURRENT APPLICATION NUMBER: US/10/367,658
CURRENT FILING DATE: 2003-02-14
PRIOR PILING DATE: 2001-02-27
PRIOR PELICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1996-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee, H.
APPLICANT: Hoe, Mee, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
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                                                                                                                            FEATURE: OTHER INFORMATION: Synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: synthetic peptide US-10-367-658-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18, Application US/10367658
Publication No. US20040071724A1
GENERAL INFORMATION:
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Publication No. US20040071724A1
GENERAL INFORMATION:
                                             TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich APPLICANT: Hoe, Mee H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100..
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HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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                LENGTH: 20
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; SEQ ID NO 12
; LENGTH: 20
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FRATURE:
; PRATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-668-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
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US-10-367-668-14
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Sequence 12, Application US/10367668

Publication No. US20040071725A1

GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Usaylew, Mark

TITLE OF INVERTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461072

CUNRENT APPLICATION NUMBER: US/10/367,668

CURRENT FILING DATE: 1998-02-13

PRIOR APPLICATION NUMBER: US 09/794,517

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, wee H.
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Wark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461072
CURRENT APPLICATION NUMBER: US/10/367,668
CURRENT FILING DATE: 2003-02-14
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR PILING DATE: 2001-02-27
PRIOR PILING DATE: 2001-02-27
PRIOR PILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) OTHER INFORMATION: Bynchetic peptide US-10-367-668-11
                                                                                                        Sequence 11, Application US/10367668
Publication No. US20040071725A1
GENERAL INFORMATION:
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1 HWDFAWPW 8
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APPLICANT: Rothman, James B.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hack, Mee, H.
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERRICE: 11746/461072
CURRENT APPLICATION NUMBER: US/10/367,668
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US 09/794,517
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1996-08-16
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NOS: 349
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotheraples
                                                                 Gaps
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0. Thath
Query Match
100.0%; Score 64, DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                 Sequence 13, Application US/10367668
Publication No. US20040071725A1
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Publication No. US20040071725A1
GENERAL INFORMATION:
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GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Hoe, Mee H.

APPLICANT: Houghton, Alan

APPLICANT: Maybew, Mark

ITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REPERENCE: 11746/461072

CURRENT PILING DATE: 2003-02-14

PRIOR FILING DATE: 2001-02-7

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR FILING DATE: 1996-08-16

PRIOR PILING DATE: 1996-08-16

PRIOR PLING DATE: 1995-08-18

PRIOR PLING DATE: 1995-08-18

PRIOR PLING DATE: 1995-08-18

PRIOR PLING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maynew, Mark
APPLICANT: Maynew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461072
CURRENT FAPLICATION NUMBER: US /01/0367,668
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,517
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR FILING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
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PRIOR FILING DATE: 1995-08-18
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PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
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, OTHER INFORMATION: synthetic peptide
US-10-367-668-16
                                                                                                                                                                Sequence 16, Application US/10367668
Publication No. US20040071725A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
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1 HWDFAWPW 8
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APPLICANT: Hartl, F. Ulrich
APPLICANT: How, Mee H.
APPLICANT: Howhton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REPERENCE: 11746/461072
CURRENT APPLICATION NUMBER: US/10/367,668
CURRENT FILING DATE: 2003-02-14
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                 CURRENT FERENCE: 1179/910/367,668
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,517
PRIOR FILING DATE: 2003-02-14
PRIOR FILING DATE: 2000-02-27
PRIOR PILING DATE: 1996-02-13
PRIOR PILING DATE: 1996-02-13
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1996-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR PLING DATE: 1995-08-18
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PRIOR PILING DATE: 2001-02-27
PRIOR PLING DATE: 2001-02-27
PRIOR PLING DATE: 1998-02-13
PRIOR PLING DATE: 1998-02-13
PRIOR PLING DATE: 1996-08-16
PRIOR PLING DATE: 1995-08-16
PRIOR PLING DATE: 1995-08-18
PRIOR PLING DATE: 1995-08-18
PRIOR PLING DATE: 1995-08-18
PRIOR PLING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SSOTUMARE: WordPerfect 8.0 for Windows
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) OTHER INFORMATION: synthetic peptide US-10-367-668-14
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US-10-367-668-15
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ORGANISM: Artificial Sequence
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FILE REFERENCE: 11746/461072
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TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies FILE REFERENCE: 11746/461072
CURRENT APPLICATION NUMBER: 02/10/367,668
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,517
PRIOR APPLICATION NUMBER: US 09/794,517
PRIOR APPLICATION NUMBER: PT/US96/13363
PRIOR PILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-16
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
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PRIOR PILING DATE: 1995-08-18
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SOFTWARE: WORDPERCECE 8:0 for Windows
SOFTWARE: WORDPERCECE 8:0 for Windows
SEQ ID NO:3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 212, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION.

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awagati, Qala

APPLICANT: Maybew, Mark

APPLICANT: Hoe, Me

TITLE OF INVENTION METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTORIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REPERRENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2000-01-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FRAESEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: synthetic peptide US-10-367-668-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 214, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION: APPLICANT: Houghcon, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens derivative
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ORGANISM: Artificial Sequence
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Matches 8: Conserve
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Best Local Similarity
Matches 8; Conserv
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US-10-258-144-212
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US-10-258-144-214
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LENGTH: 20
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APPLICANT: Hartl, F. Ulrich; APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan; APPLICANT: Takechi, Yoshizumi
         SOFTWARE: WordPerfect 8.0 for Windows SEQ ID NO 17 LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTHER INFORMATION: synthetic peptide US-10-367-668-18
                                                                                                                                                                                               ; OTHER INFORMATION: synthetic peptide US-10-367-668-17
                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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HWDFAWPW 8
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF "MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION UNDER: 60/197,642
PRIOR PLING DATE: 2000-40-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FREESE FOR Windows Version 3.0
SEG ID NO 225
LENGTH: 20
                              APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPRENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PRACEC for Windows Version 3.0
: SOFTWARE: 20
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COTHER INFORMATION: Any amino acid regidue
US-10-258-144-224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-258-144-225
; Sequence 225, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
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; Sequence 234, Application US/10258144
; Publication No. US20040101532A1
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ORGANISM: Homo sapiens derivative
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ORGANISM: Homo sapiens derivative
            Al-Awgati, Qais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 8; Conservative
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NAME/KEY: VARIANT
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Sequence 215, Application US/10258144

Publication No. US20040101532A1

Sequence 215, Application US/10258144

Publication No. US20040101532A1

APPLICANT: Houghton, Alan

APPLICANT: Livingeton, Phil

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Hoe, Mee

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: MOMBER: US/10/258,144

CURRENT PILIONG DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 2002-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FRASESEQ for Windows Version 3.0

SEQ ID NO 215

LENGTH: 20
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/17
PRIOR FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
SEQ ID NO 214
LENGTH: 20
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-215
                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens derivative
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Best Local Similarity 10u.
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Matches 8; Conservative
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1 HWDFAWPW 8
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US-10-258-144-224
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1 HWDFAWPW 8
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LENGTH: 20
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          APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: How, Mark
APPLICANT: How, Mer
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF ILLING DATE: 2002-10-17
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILLING DATE: 2000-04-17
PRIOR FILLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 235
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                              LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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COCATION: (10)...(12)

OTHER INFORMATION: Any amino acid residue
1S-10-258-144-235
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ORGANISM: Homo sapiens derivative
PEATURE:
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GENERAL INFORMATION:
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13 HWDFAWPW 20

RESULT 242

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Sequence of Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Livingston, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhaw, Mark

APPLICANT: Mayhaw, Mark

APPLICANT: Al-Awqati, Qais

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION WHOBER: US/10/258,144

CURRENT APPLICATION NUMBER: G0/197,642

PRIOR PELLING DATE: 2000-40-17

PRIOR FILING DATE: 2000-40-17

SEQ ID NO 244

LENGTH: 20

SEQ ID NO 244

LENGTH: 20
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Sequence 245, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Pabil
APPLICANT: How, Wark
APPLICANT: How, Wee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOWA
TITLE OF INVENTION: PROTEIN WEBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2000-10-17

PRIOR PLING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PEASEQ FOR Windows Version 3.0
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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COTHER INFORMATION: Any amino acid residue US-10-258-144-244
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US-10-258-144-254
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Gaps
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                                                                                                                                      APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERRING: 11746/4401
CURRENT FILING DATE: 2002-10-17
PRIOR PELING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 284
LENGTH: 20
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagari, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED INWUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEC ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
TYPE: PRI
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LOCATION: (9)...(11)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-284
                                         US-10-258-144-284
; Sequence 284, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 285, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens derivative
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                                                           APPLICANT: Hougher, Alan
APPLICANT: Hougher, Phil
APPLICANT: Hougher, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: G0/197,642
PRIOR PAPLICATION NUMBER: 60/197,642
PRIOR PAPLING DATE: 2000-40-17
NUMBER OF SEQ ID NOS: 502
SEG ID NO 254
LENGTH: 20
LENGTH: 20
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J APPLICANT: Houghton, Phil

APPLICANT: Livingston, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Hos, Me

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT FILING DATE: 2002-10-17

FRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SEQ ID NO 255

LENGTH: 20

LENGTH: 20
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; LOCATION: (9)...(11)
US-10-258-144-254
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NAME/KEY: VARIANT
NAME/TON: (10)...(12)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 255, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
Sequence 254, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
Matches 8; Conserv
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   0; Mismatches
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; OTHER INFORMATION: Any amino acid residue US-10-258-144-314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: VARIANT
LOCATION: (10)...(12)
OTHER INFORMATION: Any amino acid residue
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 314
LENGTH: 20
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Best Local Similarity 100.
Matches 8; Conservative
       8; Conservative
                                                                                                                  13 HWDFAWPW 20
                                                                        1 HWDFAWPW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-258-144-315
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LENGTH: 20
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       Matches
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                                                                        RESULT 248

US-10-258-144-294

i Sequence 294, Application US/10258144

j Publication No. US20040101532A1

j GENERAL INFORMATION:
   APPLICANT: Houghton, Alan
   APPLICANT: Livingston, Phil
   APPLICANT: Al-Awqati, Qais
   APPLICANT: Hoe, Mark
   APPLICANT: Hoe, Mark
   TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
   TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
   TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
   TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
   TITLE OF INVENTION NUMBER: US/10/258,144
   CURRENT FILING DATE: 2000-04-17
   PRIOR APPLICATION NUMBER: 60/197,642
   PRIOR PILING DATE: 2000-04-17
   NUMBER OF SEQ ID NOS: 502
   SOFTWARE: FRAESEQ for Windows Version 3.0
   SEQ ID NO 294
   TEACH OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF THE PART OF
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Phil

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION WINDER: US/10/258,144

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 295

LENGTH: 20
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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CTHER INFORMATION: Any amino acid residue
US-10-258-144-294
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LOCATION: (10)...(12)

OTHER INFORMATION: Any amino acid residue
US-10-258-144-295
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ORGANISM: Homo Bapiens derivative
FEATURE:
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13 HWDFAWPW 20
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Best Local Similarity
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| Sequence 314, Application US/10258144
| Sequence 314, Application US/10258144
| Publication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Hos, Mee
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| CURRENT PEPLICATION NUMBER: 105/10/258,144
| CURRENT FILING DATE: 2002-10-17
| PRIOR FILING DATE: 2000-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: MORPHER: 10746/46101
CURRENT FILLING NATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500-04-17
NUMBER OF SEQ ID NOS: 500-04-17
SOFTWARE: FESTSED for Windows Version 3.0
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100.0%; Pred. No. 0.071;
/ative 0; Mismatches 0; Indels
   Indels
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APPLICANT: Livingston, Argui,
APPLICANT: Livingston, Mark
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Aby Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/44610
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION UNBER: 60/197,642
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 384
LENGTH: 20
LENGTH: 20
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: How, Mea
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION NUMBER: 105/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2002-10-17
PRIOR PLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500
SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                              100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (9)...(11)
COTHER INFORMATION: Any amino acid residue
US-10-258-144-384
; LOCATION: (10)...(12)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-325
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 384, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
APPLICANT: Houghton, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 385, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
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ORGANISM: Homo sapiens derivative
                                                                                                                 Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                     13 HWDFAWPW 20
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LENGTH: 20
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US-10-258-144-325

JOS-10-258-144-325

Sequence 325, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, We
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT PILITE OF INVENTION NUMBER: US/10/258,144

CURRENT PILING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER: OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 325

LENGTH: 20

"WINDER TILING DATE: 2000-04-17

NUMBER: DATE: 2001-04-17

NUMBER: PASESEQ for Windows Version 3.0

SEQ ID NO 325
                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INCUCRALION:
GENERAL INCUCRALION:
APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION WHER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-40-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: FastSEQ for Windows Version 3.0
SEQ ID NO 324
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens derivative
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                          Query Match

100.0%; Score 64; DB 4; Length 20;

Best Local Similarity 100.0%; Pred. No. 0.071;

Matches 8; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                        Sequence 324, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens derivative
FEATURE:
NAME/KEY: VARIANT
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                                                                                                                                         1 HWDFAWPW 8
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Gaps

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FEATURE:

NAME/KEY: VARIANT

LOCATION: (10)...(12)

OTHER INFORMATION: Any amino acid residue
US-10-258-144-415
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NAME/KEY: VARIANT

LOCATION: (9)...(11)

OTHER INFORMATION: Any amino acid residue
US-10-258-144-424
SEQ ID NO 415
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens derivative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagati, Qais
APPLICANT: Al-Awagati, Qais
APPLICANT: Al-Awagati, Qais
APPLICANT: Al-Awagati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PACTOR MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERRINGE: 11746/46401
CURRENT FILING DATE: 2000-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
TYPE: PRI
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| Publication No. US20040101532A1
| GENERAL INFORMATION, Alan
| APPLICANT: Houghton, Alan
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Hoe, Mee
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION WUMBER: 60/197,642
| PRIOR APPLICATION NUMBER: 60/197,642
| PRIOR PELLING DATE: 2000-10-17
| MUMBER OF SEQ ID NOS: 500
| NUMBER OF SEQ ID NOS: 500
| NUMBER OF SEQ ID NOS: 500
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 64; DB 4; Length 20; 100.0%; Pred. No. 0.071; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                           // LOCATION: (10)...(12)
// OTHER INFORMATION: Any amino acid residue
US-10-258-144-385
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 414, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
  ORGANISM: Homo sapiens derivative
FEATURE:
NAME/KEY: VARIANT
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Matches 8; Conservative
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US-10-258-144-415
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i Sequence 424, Application US/10258144

j Publication No. US20040101532A1

j REDICALON No. US20040101532A1

j REDICANT: Houghton, Alan

j APPLICANT: Al-Awgari, Qais

j APPLICANT: Al-Awgari, Qais

j APPLICANT: Al-Awgari, Qais

j APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

j TITLE OF INVENTION: PROTHEDIN MEDIATED IMMUNOTHERAPY OF MELANOWA

TITLE OF INVENTION: PROTHER: US/10/258,144

CURRENT PILING DATE: 2002-10-17

CURRENT PILING DATE: 2002-10-17

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FaetSEQ for Windows Version 3.0

SEQ ID NO 424

LENGTH: 20
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| Publication No. US20040101532A1
| GENERAL INFORMATION, Alan
| APPLICANT: Houghton, Alan
| APPLICANT: Livingston, Phil
| APPLICANT: Al-wagati, Qais
| APPLICANT: Maybew, Mark
| APPLICANT: Maybew, Mark
| APPLICANT: Maybew, Mark
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
| CURRENT PELLING DATE: 11746/46401
| CURRENT FILLING DATE: 2002-10-17
| PRIOR APPLICATION NUMBER: 60/197,642
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Query Match
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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Gaps
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Moyew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERRICE: 11746/46101
CURRENT APPLICATION NUMBER: 2002-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 464
LENGTH: 20
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 435
LENGTH: 20
                                                                                                                                                                                                                                                                                          i LOCATION: (10)...(12)
i OTHER INFORMATION: Any amino acid residue
US-10-258-144-435
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; OTHER INFORMATION: Any amino acid residue US-10-258-144-464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-258-144-464
Sequence 464, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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; Publication No. US20040101532A1
; GENERAL INFORMATION:
APPLICANT: Houghton, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens derivative
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Al-Awgati, Qais
Mayhew, Mark
Hoe, Mee
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-465
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqari, Qais
APPLICANT: Al-Awqari, Qais
APPLICANT: Al-Awqari, Qais
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PRETRICES: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 434
LENGTH: 20
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 64; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 0.071; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                         NAME/KEY: VARIANT

1 LOCATION: (10)...(12)

OTHER INFORMATION: Any amino acid residue

1S-10-258-144-425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                 NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 425
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 434, Application US/10258144; Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                             TYPE: PRT ORGANISM: Homo sapiens derivative
            PRIOR FILING DATE: 2000-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         13 HWDFAWPW 20
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US-10-258-144-435
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us-10-16/-e/4-10

Sequence 10, Application US/10367674

Publication No. US20040127684A1

GENERAL INFORMATION:

APPLICANT: Horchman, James E.

APPLICANT: Horchman, James E.

APPLICANT: Horchman, Alam

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Horchman, Alam

APPLICANT: Horchman, Alam

APPLICANT: Horchman, Alam

APPLICANT: Horchman, Alam

APPLICANT: Mayhew, Mark

ITILE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461021

CURRENT FILING DATE: 2003-02-14

FRIOR PILING DATE: 2000-04-20

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR FILING DATE: 1996-08-16

PRIOR FILING DATE: 1996-08-16

PRIOR FILING DATE: 1996-08-16

PRIOR FILING DATE: 1996-08-16

PRIOR FILING DATE: 1996-08-16

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

SEQID NOS: 389

SEQID NO 10

TANGTHEN OF SEQID NOS: 389
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                                                 APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
FILE REFERENCE: 11746/46401
CURRENT PAPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-01-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PRESENCE for Windows Version 3.0
SEQ ID NO 495
LENGTH: 20
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; NAME/KEY: VARIANT
; LOCATION: (10)...(12)
; OTHER INFORMATION: Any amino acid residue
05-10-258-144-495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PEATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-674-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                   Al-Awgati, Qais
Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |||||||
13 HWDFAWPW 20
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; Sequence 494, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Livingston, Phil
; APPLICANT: How, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; PILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
; FILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
; FILE REPERENCE: 11746/46401
; CURRENT APPLICATION NUMBER: US/10/258,144
; PRIOR FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SEQ ID NO 494
; LENGTH: 20

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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: PEASERO for Windows Version 3.0
SEQ ID NO 465
LENGTH: 20
TYPE: PRI
CREANISM: Homo sapiens derivative
FEATURE: NUMBER: MATHORITE STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT STATEMENT 
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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I LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
1S-10-258-144-494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: VARIANT
CCATION: (10)...(12)
CTHER INFORMATION: Any amino acid residue
US-10-258-144-465
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APFLICANT: Houghton, Alan
APFLICANT: Takechi, Yoshizumi
APFLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
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CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 10/170,738
PRIOR PILING DATE: 2002-06-13
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 1996-04-20
PRIOR PILING DATE: 1996-04-20
PRIOR PILING DATE: 1996-08-16
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PRIOR PILING DATE: 1995-08-18
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PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 12
LENGTH: 20
                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: synthetic peptide US-10-367-674-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetic peptide US-10-367-674-13
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Publication No. US20040127684A1
                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
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13 HWDFAWPW 20
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                    FEATURE:
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4610211
CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT PILING DATE: 2003-02-14
PRIOR FILING DATE: 2003-06-13
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR PILING DATE: 1998-02-13
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APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mar.
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4610211
CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT FILING DATE: 2003-0214
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PRIOR APPLICATION NUMBER: US 10/170,738

PRIOR FILING DATE: 2002-06-13

PRIOR PILING DATE: 2000-04-20

PRIOR PILING DATE: 2000-04-20

PRIOR PILING DATE: 1998-02-13

PRIOR PILING DATE: 1998-02-13

PRIOR PILING DATE: 1996-08-16

PRIOR PILING DATE: 1996-08-16

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

PRIOR PILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

SOFTWARE: WordPerfect 8.0 for Windows

SEQ ID NO 11
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; OTHER INFORMATION: synthetic peptide
US-10-367-674-11
                                                                                                                                             US-10-367-674-11
; Sequence 11, Application US/10367674
; Publication No. US20040127684A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                             APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich APPLICANT: Hoe, Mee H. APPLICANT: Houghton, Alan
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Best Local Similarity 100.
Matches 8; Conservative
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HWDFAWPW 20
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Gaps

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RESULT 273
US-10-367-674-17
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Sequence 15, Application US/10367674

Publication No. US20040127684A1

GENERAL INFORMATION:

APPLICANT: Hartl, F. UIrich

APPLICANT: Hartl, F. UIrich

APPLICANT: Hartl, F. UIrich

APPLICANT: Hartl, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Mark

TITLE OF INVERTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/4610211

CURRENT APPLICATION NUMBER: US/10/367,674

CURRENT APPLICATION NUMBER: US 09/011,645

PRIOR APPLICATION NUMBER: US 09/552,868

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

SOFTWARE: WordPerfect 8:0 for Windows

SEQ ID NO 15
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                                                                        APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Hoat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4610211
CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT FILING DATE: 2003-02-14
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                      | CURKENI FILING DATE: 2002-02-14|
| PRIOR PILING DATE: 2002-06-13|
| PRIOR APPLICATION NUMBER: US 09/552,868|
| PRIOR APPLICATION NUMBER: US 09/011,645|
| PRIOR PILING DATE: 2000-04-20|
| PRIOR PILING DATE: 1998-02-13|
| PRIOR FILING DATE: 1998-08-16|
| PRIOR PILING DATE: 1996-08-16|
| PRIOR PILING DATE: 1995-08-18|
| PRIOR FILING DATE: 1995-08-18|
| PRIOR FILING DATE: 1995-08-18|
| PRIOR FILING DATE: 1995-08-18|
| NUMBER OF SEQ ID NOS: 349|
| SEQ ID NO 14|
| LENGTH: 20|
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; OTHER INFORMATION: synthetic peptide
US-10-367-674-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
APPLICANT: Rothman, James E. APPLICANT: Hartl, F. Ulrich APPLICANT: Hoe, Mee H.
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13 HWDFAWPW 20
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Sequence 16, Application US/10367674
; Sequence 16, Application US/10367674
; Publication No. US20040127684A1
; GENERAL INFORMATION:
; APPLICANT: Hotchman, James E.
APPLICANT: Hotchman, Alam
; APPLICANT: Houghton, Alam
; APPLICANT: Houghton, Alam
; APPLICANT: Houghton, Alam
; APPLICANT: Takechi, Yoshizum
APPLICANT: Takechi, Yoshizum
APPLICANT: Hotchman, Mark
; TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
; TITLE REFERENCE: 11146/461021
CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT PILING DATE: 2003-02-14
; PRIOR PILING DATE: 2000-06-13
; PRIOR PLING DATE: 1996-08-18
; PRIOR PILING DATE: 1996-08-18
; PRIOR PILING DATE: 1996-08-18
; PRIOR PILING DATE: 1995-08-18
; PRIOR PILING DATE: 1995-08-18
; PRIOR PILING DATE: 1995-08-18
; NUMBER OF SEQ ID NOS: 349
; SEG ID NO 16
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Publication No. US20040127684A1
GENERAL INPORMATION:
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Heat Shock Protein-Based Vaccines and Immunotherapies
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Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
FRATURE:
OTHER INFORMATION: synthetic peptide
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: synthetic peptide
US-10-367-674-15
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APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Houpton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Sohock Protein-Based Vaccines and Immunotherapies
     Gaps
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APPLICANT: Slusarewicz, Paul
APPLICANT: Slusarewicz, Paul
APPLICANT: Andjellic, Sofija
APPLICANT: Barber, Brian
TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF INVENTION: IMMUNOTHERAPIES
FILE REFERENCE: 849-405-999
CURRENT APPLICATION NUMBER: US/10/776,521B
CURRENT FILING DATE: 2004-02-12
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     Indels
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0
     Mismatches
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PRIOR FILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 1908-02-13
PRIOR PILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: PCT/US96/13363
PRIOR PILING DATE: 1995-08-16
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
LENGTH: 20
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CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT FILING DATE: 2003-02-14
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PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBED. HE ANALYST
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APPLICANT: Prince-Cohane, Kenya
APPLICANT: Mehta, Sunil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
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     8; Conservative
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Best Local Similarity
Matches 8; Conserva
                                                     1 HWDFAWPW 8
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US-10-776-521B-369
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Hartl, F. Ulrich
APPLICANT: Howey Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshlaumi
APPLICANT: Maybew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/4610211
CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT APPLICATION NUMBER: US 10/170,738
PRIOR PILING DATE: 2000-06-13
PRIOR PILING DATE: 1209-06-13
PRIOR PILING DATE: 1209-06-16
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8:0 for Windows
SEQ ID NO 18
LENGTH: 120
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                                CURRENT FILING DATE: 2003-02-14

PRIOR APPLICATION NUMBER: US 10/170,738

PRIOR FILING DATE: 2002-06-13

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 2998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-06-16

PRIOR PRILING DATE: 1996-08-16

PRIOR FILING DATE: 1995-08-16

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

SOUTWARE: WordPerfect 8.0 for Windows

SEQ ID NO 17

LENGTH: 20
                     CURRENT APPLICATION NUMBER: US/10/367,674
CURRENT FILING DATE: 2003-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: synthetic peptide US-10-367-674-17
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Publication No. US20040127684A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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FILE REFERENCE: 11746/4610211
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
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Sequence 374, Application US/10776521B

Publication No. US20050202033A1

Publication No. US2005020203A1

GENERAL INFORMATION:

APPLICANT: Fletchner, Jessica

APPLICANT: Fletchner, Jessica

APPLICANT: Sunia

APPLICANT: Andjelic, Paul

APPLICANT: Andjelic, Paul

APPLICANT: Barber, Brian

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION UNMBER: 60/403,417

PRIOR APPLICATION NUMBER: 60/463,746

PRIOR APPLICATION NUMBER: 60/463,746

PRIOR PLIING DATE: 2003-04-18

PRIOR PLIING DATE: 2003-04-18

PRIOR PLIING DATE: 2003-02-13

NUMBER OF SEQ ID NOS: 419

SOFTWARE: FREESEQ FOR WINGOWS VERSION 4.0

SEQ ID NO 374

LENGTH: 20

LENGTH: 20

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LENGTH: 20
                    APPLICANT: Slusaredicz, Paul
APPLICANT: Slusaredicz, Paul
APPLICANT: Slusaredicz, Sofija
APPLICANT: Barber, Barian
TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
TITLE OF INVENTION: IMMUNOTHERAPIES
FILE REPREBACES: 8449-405-999
CURRENT APPLICATION NUMBER: 05/503,417
PRIOR APPLICATION NUMBER: 60/503,417
PRIOR PILING DATE: 2003-09-108
PRIOR FILING DATE: 2003-04-18
PRIOR FILING DATE: 2003-04-18
PRIOR PELING DATE: 2003-04-11
PRIOR PELING DATE: 2003-04-13
PRIOR FILING DATE: 2003-04-13
PRIOR FILING DATE: 2003-04-13
SOFTWARE: FastSEQ for Windows Version 4.0
IENGTH: 20
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US-10-776-521B-374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
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ORGANISM: Artificial Seguence
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Best Local Similarity 100.(
Matches 8; Conservative
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US-10-776-521B-374
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GENERAL INFORMATION:
APPLICANT: Fletchner, Jessica
APPLICANT: Prince-Cohane, Kenya
APPLICANT: Prince-Cohane, Kenya
APPLICANT: Mahte, Sunil
APPLICANT: Slusarewicz, Paul
APPLICANT: Barber, Brian
ITILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
ITILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND
ITILE OF INVENTION: IMMUNOTHERAPIES
FILE REFERENCE: 8449-405-99
CURRENT APPLICATION NUMBER: 06/503,417
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR APPLICATION NUMBER: 60/463,746
PRIOR PRILING DATE: 2003-09-16
PRIOR PPLING DATE: 2003-04-18
PRIOR PPLING DATE: 2003-04-18
PRIOR PPLING DATE: 2003-02-13
PRIOR SPELING DATE: 2003-02-13
PRIOR SPELING DATE: 2003-02-13
NUMBER OF SEQ ID NOS: 419
: LENGTH: 20
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100.0%; Score 64; DB 5; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 5; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
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PRIOR APPLICATION NUMBER: 60/462,469
PRIOR FILING DATE: 2003-04-11
PRIOR PPLING DATE: 2003-02-13
NUMBER OF SEQ ID NOS: 419
SOFTWARE: FARESEQ for Windows Version 4.0
SEQ ID NO 369
LENGTH: 20
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Sequence 373, Application US/10776521B
Sequence 373, Application US/10776521B
Publication No. US20050202033A1
GENERAL INFORMATION:
APPLICANT: Fletchner, Jessica
APPLICANT: Prince-Cohane, Kenya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 371, Application US/10776521B Publication No. US20050202033A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Hybrid antigen US-10-776-521B-369
                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-10-17
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APPLICANT: Houghton, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
CURRENT APPLICATION NUMBER: US/10/258,144
                                                                                100.0%; Score 64; DB 4; Length 21; 100.0%; Pred. No. 0.074;
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100.0%; Score 64; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                     0; Indels
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; OTHER INFORMATION: Any amino acid residue
US-10-258-144-264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 265
LENGTH: 21
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 304
LENGTH: 21
                                                                                                                                                                                                                                                                                                                       RESULT 282
US-10-258-144-265
US-10-258-144-265

Sequence 265, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-04-17
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PEATURE:
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ORGANISM: Homo sapiens derivative
                                                                                Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-264

Sequence 264, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/4641

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 2001-017

NUMBER: OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 264

LENGTH: 21
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100.0%; Score 64; DB 5; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.071;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                      VS-10-820-067A-878

Sequence 878, Application US/10820067A

Publication No. US20050214312A1

GENERAL INFORMATION:
APPLICANT: Fletchner, J.
APPLICANT: Mehta, S.
APPLICANT: Mehta, S.
APPLICANT: Mehta, S.
APPLICANT: Mehta, S.
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APPLICANT: Mehta, S.
APPLICANTON VUMBER: US/10/820,067A

CURRENT FILING DATE: 2003-04-18

PRIOR FILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR APPLICATION NUMBER: 60/463,746

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

PRIOR PILING DATE: 2003-04-18

SEQ ID NOS: 926

SSCTWARE PRESENCE (or Windows Version 4.0)

SSQ ID NO 8'8

LENGTH: 20
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
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1 HWDFAWPW 8
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j Sequence 305, Application US/10258144

j Publication No. US20040101532A1

geneRAL INFORMATION:
   APPLICANT: Houghton, Alan
   APPLICANT: Houghton, Phil
   APPLICANT: How, Mark
   APPLICANT: How, Mark
   APPLICANT: How, Mark
   APPLICANT: How, Mark
   APPLICANT: MAYNAW, MATHODS AND COMPOSITIONS FOR HEAT SHOCK
   TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
   FILE REFERENCE: 11746/46401
   CURRENT PILING DATE: 2000-04-17
   PRIOR APPLICATION NUMBER: 60/197,642
   PRIOR PILING DATE: 2000-04-17
   NUMBER OF SEQ ID NOS: 502
   SOFTWARE: FastSEQ for Windows Version 3.0
   SEQ ID NO 305
   LENGTH: 21
   TYPE: DOS: 502
   TYPE: DOS: 502
   TYPE: DOS: 502
   SEQ ID NO 305
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Sequence 334, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Al-Awqari, Qais

APPLICANT: How, Mark

APPLICANT: How, Mark

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT PILING DATE: 2002-10-17

PRIOR PILING DATE: 2000-04-17
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                                                                                                                                             Query Match 100.0%; Score 64; DB 4; Length 21; Best Local Similarity 100.0%; Pred. No. 0.074; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 64; DB 4; Length 21; Best Local Similarity 100.0%; Pred. No. 0.074; Matches 8; Conservative 0; Mismatches 0; Indels
; FEATURE:
; NAME/KEY: VARIANT
; LOCATION: (9)...(11)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-304
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CCATION: (11)...(13)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-305
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SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 334
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERBNCE: 11/146/46401
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PAPLICATION NUMBER: 60/197,642
PRIOR PAPLICATION STER 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PSECSEQ for Windows Version 3.0
SEQ ID NO 355
LENGTH: 21
TYPE: PRT
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Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqeti, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
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                                                                                                                                                                                                                                                                 ; LOCATION: (9)...(11)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-354
                      PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 354
LENGTH: 21
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; Sequence 364, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens derivative
  CURRENT FILING DATE: 2002-10-17
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Matches 8; Conservative
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
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; Sequence 354, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Phil
; APPLICANT: APPLICANT: APPLICANT: APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; TILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
                                                                                                                                                                                                                                                                                       100.0%; Score 64; DB 4; Length 21; 100.0%; Pred. No. 0.074; Artive 0; Mismatches 0; Indels
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                                                                                                                                                                  NAME/KEY: VARIANT

CCATION: (9)...(11)

G THER INFORMATION: Any amino acid residue
US-10-258-1444-344
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NAME/KEY: VARIANT
LOCATION: (11)...(13)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-345
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 344
LENGTH: 21
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CURRENT APPLICATION NUMBER: US/10/259,144
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Publication No. US20040101532A1
GENERAL INFORMATION:
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                                                                                                                     ORGANISM: Homo sapiens derivative
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Best Local Similarity luv.
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Best Local Similarity 100.
Matches 8; Conservative
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LENGTH: 21
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Fublication No. US2004101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Al-Awgati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOWA

TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOWA

FILE REFERENCE: 11746/46401

CURRENT FILING DATE: 2002-10-17

PRIOR FILING DATE: 2002-01-17

PRIOR PILICATION NUMBER: 60/197,642

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SEPTIMES FERENCE: 110-04-17

NUMBER OF SEQ ID NOS: 502

SEPTIMES FARESEQ for Windows Version 3.0
                      TITLE OF INVENTION: RETHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILL SOURCE : 2002-10-17
CURRENT PAPLICATION NUMBER: 60/197,642
PRIOR PRICH DATE: 2000-04-17
NUMBER OF ERQ ID NOS: 502
SOFTWARE: FastSRQ for Windows Version 3.0
SEQ ID NO 374
LENGTH: 21
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NAME/KEY: VARIANT
LOCATION: (11)...(13)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Homo sapiens derivative
FEATURE:
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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US-10-258-144-394
Sequence 394, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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Matches 8; Conservative
         Mayhew, Mark
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LENGTH: 21
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APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/44401
CURRENT APPLICATION NUMBER: 05/197,642
PRIOR PILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
LENGTH: 21
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TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 364
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100.0%; Score 64; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0: Tndals
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COTHER INFORMATION: Any amino acid residue
US-10-258-144-364
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COTHER INFORMATION: Any amino acid residue
US-10-258-144-365
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US-10-256-144-374
Sequence 374, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Al-Awqati, Qais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 365, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens derivative
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NAME/KEY: VARIANT
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Application US/10258144
5. US20040101532A1
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ORGANISM: Homo sapiens derivative
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ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.0
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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NAME/KEY: VARIANT
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US-10-258-144-405
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hos, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 395
LENGTH: 21
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Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
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CCATION: (9)...(11)
US-10-258-144-394
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1 LOCATION: (11)...(13)
2 OTHER INFORMATION: Any amino acid residue
US-10-258-144-395
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Homo sapiens derivative
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RESULT 297 US-10-258-144-404

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Sequence 405, Application US/10258144

Publication No. US20040101532A1

Publication No. US20040101532A1

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

APPLICANT: Hoe, Mee

ITTLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

ITTLE OF INVENTION: PROTEIN: US/10/258,144

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT PILING DATE: 2002-10-17

PRIOR FILING DATE: 2000-04-17

NUMBER: OF SEQ ID NOS: 502

SOFTWARE: PSSEQ for Mindows Version 3.0

SEQ ID NO 405

LENGTH: L.
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awdati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11466/46101
CURRENT APPLICATION NUMBER: 2002-10-17
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 404
LENGTH: 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COCATION: (9)...(11)
COTHER INFORMATION: Any amino acid residue
US-10-258-144-404
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14 HWDFAWPW 21
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US-10-286-144-445

Sequence 445, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
FAPPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
FILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
FRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502

SEQ ID NO 45
LENGTH: 21

TYPE: FREE
                                                    Sequence 444, Application US/10258144
; Sequence 444, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION; Alan
; APPLICANT: Houghton, Alan
; APPLICANT: Houghton, Phil
   APPLICANT: MayNew, Mark
; APPLICANT: MayNew, Mark
; APPLICANT: MayNew, Mark
; APPLICANT: Ho, Me
   TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION NUMBER: US/10/258,144
; CURRENT PELICATION NUMBER: US/10/258,144
; CURRENT PLING DATE: 2002-10-17
; PRIOR APPLICATION NUMBER: 60/197,642
; PRIOR APPLICATION NUMBER: 200-04-17
; NUMBER OF SEQ ID NOS: 502
; SEQ ID NO 444
LENGTH: 21
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100.0%; Score 64; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
100.0%; Score 64; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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FEATURE:
NAME/KEY: VARIANT
OCCATION: (11)...(13)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens derivative
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1 HWDFAWPW

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Sequence 454 Application US/1028144

Publication 454 Application US/1028144

Publication No. USZODOGOUSSZAA

GENERAL WFORWAIDN:

APPLICANT: LAVINGENCO. PHIL

APPLICANT: LAVINGENCO. PHIL

APPLICANT: HORNATION: PHIL

APPLICANT: HORNATION: PHIL

APPLICANT: MAPAN. MARK

APPLICANT: MAPAN. MARK

TITIES OF INMERIOR DATE: 2000-04-17

FILE SEGRED OF DATE: 2000-04-17

WINNERS OF SEG ID NOS: 502

SEG ID NOS: 502

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SEG ID NOS: 502

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SEG ID NOS: 502

SEG ID NOS: 503

SEG ID NOS: 503

SEG ID NOS: 503

SEG ID NOS: 504

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SEG ID N
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14 HWDFAWPW 21
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; Sequence 475. Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION;
APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
APPLICANT: Maybew, Mark
; APPLICANT: Maybew, Mark
; APPLICANT: Maybew, Mark
; APPLICANT: Hoo, Mee
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA; FILE REPERBUGE: 11746/46401
; CURRENT APPLICATION NUMBER: 2002-10-17
; PRIOR PLING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFFWARE: FEBELSEQ for Windows Version 3.0
; SEQ ID NO 475
; LENGTH: 21
                                                                                                                                                                                                                                                  APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Aqqati, Qais
APPLICANT: Al-Aqqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 17746/46401
CURRENT APPLICATION NUMBER: 60/197,642
FRIOR FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 474
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens derivative
FEATURE:
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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FEATURE:
NAME/KEY: VARIANT
LOCATION: (11)...(13)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-475
                                                                                                                                                                                      Sequence 474, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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Matches 8, Conservative
  8; Conservative
                                                                          14 HWDFAWPW 21
                                            1 HWDFAWPW 8
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US-10-258-144-475
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    Matches
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NUMBER OF SEQ ID NOS: 502
SOFTHARE: FastSEQ for Windows Version 3.0
SEQ ID NO 279
LENGTH: 24
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-258-144-283
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LENGTH: 25
TYPE: PRT
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Sequence 378, Application US/10776521B

Publication No. US20050202033A1

GENERAL INFORMATION:

APPLICANT: Fletchner, Jessica

APPLICANT: Fletchner, Jessica

APPLICANT: Application No. Setija

APPLICANT: Application Solija

APPLICANT: Application Solija

APPLICANT: Application Solija

APPLICANT: Application Solija

APPLICANT: Application Solija

APPLICANT: Barber, Brian

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

TITLE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION: IMPROVED HEAT SHOCK PROTEIN-BASED VACCINES AND

FILE OF INVENTION NUMBER: 60/462, 469

FRIOR FILING DATE: 2003-09-11

FRIOR FILING DATE: 2003-09-11

FRIOR FILING DATE: 2003-09-11

FRIOR FILING DATE: 2003-09-11

FRIOR FILING DATE: 2003-09-11

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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
APPLICANT: How, Mex
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR APPLICATION NUMBER: 0202-10-17
PRIOR APPLICATION NUMBER: 0202-10-17
PRIOR APPLICATION NUMBER: 0202-10-17
PRIOR PLING DATE: 2000-04-17
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100.0%; Score 64; DB 5; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.074;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                           Query Match 100.0%; Score 64; DB 4; Length 21; Best Local Similarity 100.0%; Pred. No. 0.074; Matches 8; Conservative 0; Mismatches 0; Indels
       ; LOCATION: (11)...(13)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-485
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Hybrid antigen US-10-776-521B-378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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US-10-776-521B-378
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Gaps
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPERRENCE: 11746/4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-04-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 283
LENGTH: 24
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Markw, Mark
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
FRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PABLESEQ for Windows Version 3.0
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                                                                                                                  100.0%; Score 64; DB 4; Length 24; 100.0%; Pred. No. 0.082;
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100.0%; Pred. No. 0.082;
ative 0; Mismatches 0; Indels
                                                                                                                                                                          0; Indels
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 283, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
; TYPE: PRT; ORGANISM: Homo sapiens derivative US-10-258-144-279
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Gaps
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  Gape
                                                                                                                                                                                                                                                                                           APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagari, Qais
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 2002-01-7
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PESESSEQ for Windows Version 3.0
SETWARE: 25
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100.0%; Score 64; DB 4; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
  Mismatches
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 229
LENGTH: 25
TYPE: PRT
                                                                                                                                                                                                                   Sequence 223, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
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; Sequence 229, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
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Best Local Similarity 100.
Matches 8; Conservative
  8; Conservative
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 0S/197,642
PRIOR FILING DATE: 2002-10-17
PRIOR FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 219
LENGTH: 25
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0
                                                                     100.0%; Score 64; DB 4; Length 25;
100.0%; Pred. No. 0.085;
ative 0; Mismatches 0; Indels
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. Sequence 219, Application US/10258144
. Publication No. US20040101532A1
. GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                               Sequence 211, Application US/10258144
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; ORGANISM: Homo sapiens derivative
US-10-258-144-219
; ORGANISM: Homo sapiens derivative US-10-258-144-210
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                                                                   Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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HWDFAWPW 8
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Best Local Similarity
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Publication No. US20040101532A1
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US-10-258-144-253
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US-10-239

Sequence 239, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alm
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: WarHODS AND COMPOSITIONS FOR HEAT SHOCK
ITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FRAESEQ for Windows Version 3.0

SEQ ID NO 239

LENGTH: 25

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APPLICANT: Livingston, Phil

APPLICANT: Livingston, Oais

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

TITLE OF INVERTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVERTION: PROTEIN MEDIATED INVUNOTHERAPY OF MELANOWA

TITLE OF INVERTION: PROTEIN MEDIATED INVUNOTHERAPY OF MELANOWA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SEQ ID NO 233

LENGTH: 25

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LENGTH: 25
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100.0%; Score 64; DB 4; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                      Sequence 233, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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US-10-258-144-243
; Sequence 243, Application US/10258144
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Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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GENERAL INFORMATION, Alan

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mermon Mark

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REPERENCE: 11746/446401

CURRENT APPLICATION NUMBER: US/10/258,144

FRIOR APPLICATION NUMBER: 0012-10-17

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOC ID NOS: 502

LENGTH: 25

LENGTH: 25
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US-10-258-144-263
i Sequence 253, Application US/10258144
j Publication No. US20040101532A1
j GENERAL INFORMATION:
i APPLICANT: Houghton, Alan
j APPLICANT: Livingston, Phil
j APPLICANT: Al-Awgati, Qais
j APPLICANT: Hoe, Mee
j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
j TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
j FILE REFERENCE: 11746/446101
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APPLICANT: He, we merupos and compositions for heat shock tille of invention: Protein mediated immunorherapy of melanoma file of invention: Protein mediated immunorherapy of melanoma file of invention: 11746/46401 current application number: us/10/258,144 current filing date: 2002-10-17 current filing date: 2002-10-17 prior Filing date: 2000-04-17 number of seq id nos: 502 prior seq id nos: 502 seq id nos: 502 seq id no 253 iength: 25
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Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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; Sequence 259, Application US/10258144
; Publication No. US20040101532A1
                                                                                                                                                                                                                                                                                                              ) ORGANISM: Homo sapiens derivative US-10-258-144-253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-259
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Matches 8; Conservative
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CURRENT APPLICATION NUMBER: 18/10/258,144

PRIOR PLING DATE: 2002-10-17

PRIOR PLING DATE: 2002-10-17

PRIOR PLING DATE: 2002-0-17

PRIOR PLING DATE: 2002-0-41

PRIOR PLING DATE: 2002-0-41

PROPER PLING DATE: 2000-0-41

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Sequence 319, Application No. US20040101532A1

GENERAL INFORMATION.

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Livingston, Phil

APPLICANT: How, Mark

APPLICANT: How, Mark

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA

CURRENT APPLICATION NUMBER: 108/10/258,144

CURRENT PEPLICATION NUMBER: 60/197,642

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FRELSE (60/197,642)

SOFTWARE: FRELSE (60/197,642)

SOFTWARE: FRELSE (60/197,642)

SOFTWARE: FRELSE (60/197,642)

TYPE: PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NO 319

LENGTH: 25

TYPE: PRIOR FILING DATE: 2000-04-17

SOFTWARE: PRIOR FILING DATE: 2000-04-17

SOFTWARE: PRIOR FILING DATE: 2000-04-17

SOFTWARE: PRIOR FILING DATE: 2000-04-17

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100.0%; Score 64; DB 4; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                               100.0%; Score 64; DB 4; Length 25; 100.0%; Pred. No. 0.085;
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                                                                                                    Query Match
Best Local Similarity
Matches 8; Conserv
     US-10-258-144-303
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US-10-258-144-299
Sequence 299, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingeron, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
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APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: MATHORS AND COMPOSITIONS FOR HEAT SHOCK
ITLE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 108/10/258,144
CURRENT PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 299
LENGTH: 25
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US-10-228-144-303
i Sequence 303, Application US/10258144
j Publication No. US20040101532A1
j GENERAL INFORMATION:
   APPLICANT: Houghton, Alan
j APPLICANT: Al-Awqati, Qais
   APPLICANT: Mayhaw, Mark
   APPLICANT: Al-Awqati, Qais
   TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
   TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
   TITLE OF ILLING DATE: 2002-10-17
   CURRENT APPLICATION NUMBER: US/10/258,144
   CURRENT APPLICATION NUMBER: 60/197,642
   PRIOR FILING DATE: 2000-04-17
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Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 293 LENGTH: 25
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 303
LENGTH: 25
                                                                                                                                               ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-293
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ORGANISM: Homo sapiens derivative
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                                                                                                                                                               APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11146/46101
CURRENT APPLICATION NUMBER: 2002-01-17
PRIOR PLIING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 389
LENGTH: 25
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APPLICANT: ALLAWGATI, Cais
APPLICANT: ALLAWGATI, Cais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 00/19/642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 393
LENGTH: 25
TYPE: PRT
TYPE: PRT
TYPE: PRT
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                                                                  Sequence 389, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
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; Sequence 419, Application US/10258144
; Publication No. US20040101532A1
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; ORGANISM: Homo sapiens derivative
US-10-258-144-389
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Best Local Similarity 100.
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                                                                                                                                                                                                                                     Sequence 329, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMAT: Worder Alan

APPLICANT: Livingsten, Phil

APPLICANT: Livingsten, Phil

APPLICANT: How, Mark

APPLICANT: How, Mer

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: MOMBER: US/10/258,144

CURRENT APPLICATION NUMBER: 05/10/10/258,144

CURRENT FILING DATE: 2000-10-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOSS: 500

SOFTWARE: FREEKEQ FOR WINDOWS VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Howardi, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERBUCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICANT NUMBER: 60/197,642
PRIOR PELING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: PRACEEQ for Windows Version 3.0
SEQ ID NO 333
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 333, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-329
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Best Local Similarity الاست
الا الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد الاستخداد ا
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Best Local Similarity 100.
Matches 8; Conservative
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LENGTH: 25
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US-10-258-144-433

Sequence 433, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

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APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

FILE REFERENCE: 11746/46401

CURRENT PILING DATE: 2000-10-17

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PattSEQ for Windows Version 3.0

SEQ ID NO 433

LENTH: AL-AWAGE

CURRENT: AL-AWAGE

CORGANISM: Home sapiens derivative

US-10-258-144-433
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Sequence 439, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
CURRENT APPLICATION NUMBER: US/10/258,144
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APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PASTESEQ FOR WINDOWS VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 64, DB 4; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Homo sapiens derivative
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Best Local Similarity luv...
And By Conservative
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LENGTH: 25
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| Sequence 423, Application US/10258144
| Publication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Al-Awqati, Qais
| APPLICANT: Mayhew, Mark
| APPLICANT: Mayhew, Mark
| APPLICANT: Al-Awqati, Qais
| APPLICANT: Al-Awqati, Qais
| APPLICANT: Houghton: PROTEIN Phil
| APPLICANT: How, Mark
| TITLE OF INVENTION: PROTEIN WEDIATED INFUNDITHERAPY OF MELANOMA
| TITLE OF INVENTION: PROTEIN WEDIATED OF UNCHERAPY OF MELANOMA
| TITLE OF ILLY GAIS
| CURRENT FILING DATE: 2002-10-17
| PRIOR FILING DATE: 2000-04-17
| NUMBER OF SEQ ID NOS: 502
| SEQ ID NO 423
| LENGTH: 25
                   APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF ILLING DATE: 2002-10-17
CURRENT PILLING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOUTHARE: PASSEQ for Windows Version 3.0
SEQ ID NO 419
LENGTH: 25
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100.0%; Pred. No. 0.085;
vative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 64; DB 4; Length 25; Best Local Similarity 100.0%; Pred. No. 0.085; Matches 9; Conservative 0; Mismatches 0; Indels
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; Sequence 429, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INPORMATION;
APPLICANT: Houghton, Alan
; APPLICANT: Al-Angati, Oais
; APPLICANT: Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ) ORGANISM: Homo sapiens derivative US-10-258-144-423
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDFAWPW 8
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     GENERAL INFORMATION:
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APPLICANT: Livingston, Pain.
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
APPLICANT: Mayhaw, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTUSIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/4641
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR PILING DATE: 2000-46-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PESECRE for Windows Version 3.0
SEQ ID NO 499
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Hougheati, Qaia
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILLING DATE: 2000-210-17
PRIOR PLLING DATE: 2000-210-17
PRIOR PLLING DATE: 2000-04-17
NUMBER OF SEC ID NOS: 502
SOFTWARE: FastSEC for Windows Version 3.0
SOFTWARE: 25
TYPE: PRT
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                                                                                                                                                    100.0%; Score 64; DB 4; Length 25; 100.0%; Pred. No. 0.085; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 473, Application US/10258144; Publication No. US20040101532A1; GENERAL INFORMATION:
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                               ; LENGTH: 25
; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-469
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                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-258-144-499
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       SEQ ID NO 469
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; Sequence 443, Application No. US20040101532A1
; Beblication No. US20040101532A1
; GENERAL INFORMATION:
    APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Al-Awdati, Qais
; APPLICANT: Maybew, Mark
; TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: DATE: 2002-10-17
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT FILING DATE: 2002-10-17
; PRIOR FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: PRESEQ for Windows Version 3.0
; SEQ ID NO 443
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT APPLICATION NUMBER: 61010/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500-04-17
NUMBER OF SEQ ID NOS: 500-04-17
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100.0%; Score 64; DB 4; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FRSUSEQ for Windows Version 3.0
SEQ ID NO 439
LENGTH: 25
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-439
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; ORGANISM: Homo sapiens derivative
US-10-258-144-443
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1 HWDFAWPW 8
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HWDFAWPW 8
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Sequence 309, Application US/10258144

Publication No. US2040101532A1

GENERAL INFORMATION,
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
APPLICANT: Hoe, Mee
APPLICANT: Hoe, Mee
CINTERION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502

SOFTWARE FREEESE for Windows Version 3.0
SOFTWARE FREEESE for Windows Version 3.0
LENGTH: 26
TUNDED TO NO 309
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Sequence 313, Application US/10258144

Sequence 313, Application US/10258144

Publication No. US20040101532A1

APPLICANT: Houghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Al-Awgati, Cais

APPLICANT: Hoe, Mee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT PILLING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR FILLING DATE: 2002-10-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PastSEQ for Windows Version 3.0

SEQ ID NO 313

LENGTH: 26

TYPE: PRT

ORGANISM: Homo sapiens derivative

US-10-258-144-313
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100.0%; Pred. No. 0.088;
/ative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 26;
Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 8; Conservative
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HWDFAWPW
                                  1 HWDFAWPW
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US-10-258-144-309
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US-10-258-144-273
is Sequence 273, Application US/10258144
j Publication No. US20040101532A1
is GENERAL INFORMATION:
j APPLICANT: Hodehton, Alan
j APPLICANT: Livingston, Phil
j APPLICANT: Hadehwe, Mark
j APPLICANT: Mayhew, Mark
j APPLICANT: Mayhew, Mark
j APPLICANT: Mayhew, Mark
j APPLICANT: Mayhew, Mark
j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
j TITLE OF INVENTION WINBER: US/10/258,144
j CURRENT APPLICATION NUMBER: 2002-10-17
j PRIOR APPLICATION NUMBER: 2000-04-17
j PRIOR FILING DATE: 2000-04-17
j NUMBER: PRACESEQ for Windows Version 3.0
j SEQ ID NO 273
j LENGTH: 26
                                                                                                                                                                                                                                                                                                                 Sequence 269, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Livingscon, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA
TITLE OF INVENTION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 269
LENGTH: 26
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                      Query Match
Best Local Similarity 100.0%; Pred. No. 0.085;
Matches 8; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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; ORGANISM: Homo sapiens derivative
US-10-258-144-273
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US-10-258-144-269
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                                                                                                         APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITION FOR HEAD
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 349
LENGTH: 26
TYPE: PRT
CREANISM: Homo sapiens derivative
US-10-258-144-349
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FUBLICATION NO. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Haveghton, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

APPLICANT: Hos, We

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/4440

CURRENT PILICATION NUMBER: US/10/258,144

CURRENT PILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 353

LENGTH: 26
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: A-Magati, Qais
APPLICANT: Hoe, Mee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                      Livingston, Phil
Al-Awgati, Qais
Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 8, Conservative
                                                Houghton, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW 8
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US-10-258-144-359
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RESULT 345
US-10-258-144-339
is Sequence 339, Application US/10258144
ibulication No. US20040101532A1
is GRNERAL INFORMATION:
APPLICANT: Livingston, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
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APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
CURRENT PILING DATE: 2002-10-17
FRIOR PILING DATE: 2002-10-17
FRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 339
LENGTH: 26
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| Sequence 343, Application US/10258144
| Publication No. US20040101532A1
| GENERAL INFORMATION:
| APPLICANT: Houghton, Alan
| APPLICANT: Houghton, Phil
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Al-Awgati, Qais
| APPLICANT: Al-Awgati, Qais
| APPLICANT: However, Mark
| TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR HEAT SHOCK
| TITLE OF INVENTION: PROFIEIN MEDIATED INMUNOTHERAPY OF MELANOMA
| TITLE OF INVENTION: PROFIEIN WEDIATED INMUNOTHERAPY OF MELANOMA
| TITLE OF INVENTION NUMBER: US/10/258,144
| CURRENT APPLICATION NUMBER: 60/197,642
| PRIOR FILING DATE: 2002-10-17
| WUMBER OF SEQ ID NOS: 502
| SOFTWARE: PRASESEQ for Windows Version 3.0
| SEQ ID NO 343
| LENGTH: 26
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100.0%; Score 64; DB 4; Length 26;
Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 26;
Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens derivative US-10-258-144-339
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RESULT 347

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; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-369
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1 HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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US-10-258-144-379
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Ani

APPLICANT: Houghton, Phil

APPLICANT: How, Mark

APPLICANT: Maybew, Mark

APPLICANT: Maybew, Mark

APPLICANT: Maybew, Mark

TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: 00/10/258,144

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR PRING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: PRESEQ for Windows Version 3.0

SEQ ID NO 363

LENGTH: 26
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US-10-258-144-369

i Sequence 369, Application US/10258144

i Publication No. US20040101532A1

i GENERAL INFORMATION:

i APPLICANT: Houghton, Alan

i APPLICANT: Livingston, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Hoe, Mark

i APPLICANT: Al-Awqati, Qais

i TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA

i TILLE REPERENCE: 11746/446401

i CURRENT APPLICATION UNBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA FILE REPERENCE: 11746/46401 US/10/258,144
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR PAPLICATION NUMBER: 60/197,642
PRIOR PELING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE FARESEQ for Windows Version 3.0
SEQ ID NO 3:9
LENGTH: 26
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                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-359
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDFAWPW 8
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; bedication No. US20040101532A1
; bedication No. US20040101532A1
; bedication No. US20040101532A1
; ceneral information.
; APPLICANT: Livingston, Alan
; APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
; APPLICANT: Hoe, Mer
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; ITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA; PILE REPERBURE: 11746/46101
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT PILING DATE: 2002-10-17
; PRIOR PPLICATION NUMBER: 60/197,642
; RUMBER OF SEQ ID NOS: 502
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 379
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                                                                                                                                                                                 Query Match
100.0%; Score 64; DB 4; Length 26;
Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 369
LENGTH: 26
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Gaps
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Houghton, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Hoe, Mee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46101

CURRENT FILING DATE: 2002-10-17

PRIOR PAPLICATION NUMBER: 60/197,642

PRIOR PAPLICATION NUMBER: 60/197,642

PRIOR PAPLICATION NUMBER: 60/197,642

PRIOR PAPLICATION NUMBER: 60/197,642

SOFTHARE: PERESE FOR MINDOWS VERSION 3.0

SOFTHARE: PERESE FOR MINDOWS VERSION 3.0

SOFTHARE: PERESE FOR MINDOWS VERSION 3.0
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awdati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Wee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERROR: 11746/46401
CURRENT APPLICATION NUMBER: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTMARE: PRESENCE for Windows Version 3.0
SOFTMARE: 26
LENGTH: 26
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Query Match 100.0%; Score 64; DB 4; Length 26; Best Local Similarity 100.0%; Pred. No. 0.088; Matches 8; Conservative 0; Mismatches 0; Indels
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Publication No. US20040101532A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens derivative
US-10-258-144-403
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Best Local Similarity الاست
الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستراكة الاستركة الاستركة الاستراكة الاستراكة الاستراكة الاسترا
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; Publication No. US20040101532A1
; GENERAL INFORMATION:
    APPLICANT: Houghton; Alan
    APPLICANT: Livingston, Phil
    APPLICANT: Maybew, Mark
    APPLICANT: Maybew, Mark
    APPLICANT: Maybew, Mark
    APPLICANT: Maybew, Mark
    APPLICANT: Al-Awgati, Qais
    APPLICANT: Maybew, Mark
    APPLICANT: Al-Awgati, Qais
    APPLICANT: Al-Awgati, Qais
    APPLICANT: Al-Awgati, Qais
    APPLICANT: Al-Awgati, Qais
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    APPLICANT: Al-Awgati, Qais
    APPLICANT: Al-Awgati, Qais
    CURRENT APPLICATION NUMBER: US/10/258,144
    CURRENT APPLICATION NUMBER: Go/197,642
    PRIOR PELING DATE: 2000-04-17
    NUMBER OF SEQ ID NOS: 502
    SEQ ID NO 399
    LENGTH: 26
    LENGTH: 26
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APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqari, Qais
APPLICANT: Al-Awqari, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/46401
CURRENT PILING DATE: 2000-10-17
FRICK APPLICATION NUMBER: 60/197,642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Mindows Version 3.0
SEQ ID NO 383
LENGTH: 26
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                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 4; Length 26;
Best Local Similarity 100.0%; Pred. No. 0.088;
Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 383, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
                             ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-383
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; ORGANISM: Homo sapiens derivative
US-10-258-144-399
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDFAWPW 8
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LENGTH: 26
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1 HWDFAWPW 8

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US-10-258-144-459

Sequence 459, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingarti, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: MAYNEW, MATHON WETHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
ITILE OF INVENTION: NETHODS AND COMPOSITIONS
FILE REFERENCE: 11746/46401
CURRENT FILING DATE: 2002-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOUTHWARE: FESTESEQ FOR Windows Version 3.0
US-10-258-144-453

Sequence 455, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awdati, Gais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Me
TILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TILLE REFERENCE: 11746/46401
FILLE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR PILLIATION DATE: 2002-04-17
PRIOR PILLIATION DATE: 200-04-17

NUMBER OF SEQ ID NOS: 502

SEQ ID NO 453

LENGTH: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT; ORGANISM: Homo sapiens derivative US-10-258-144-453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 8; Conservative
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Matches 8; Conservative
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LENGTH: 26
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                                                                                                                 Sequence 413, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Houghton, Phil
; APPLICANT: Houghton, Phil
; APPLICANT: How, Mark
; APPLICANT: How, Mem
; APPLICANT: How, Mem
; APPLICANT: How, Mem
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION NUMBER: US/10/258,144
; CURRENT APPLICATION NUMBER: 2002-10-17
; PRIOR PILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: PRAESEQ for Windows Version 3.0
; SEQ ID NO 413
; LENGTH: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 449, Application US/10258144
; Sequence 449, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Maybew, Mark
; APPLICANT: Al-Awqati, Qais
; APPLICANT: Maybew, Mark
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: PROTEIN NEDIATED IMMUNOTHERAPY OF MELANOMA;
; CURRENT FILING DATE: 2002-10-17
; PRIOR FILING DATE: 2002-10-17
; PRIOR FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: PRESEQ for Windows Version 3.0
; SEQ ID NO 449
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT;
CORGANISM: Homo sapiens derivative
US-10-258-144-413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.1
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDFAWPW
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: How, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: 17346/46401

CURRENT APPLICATION NUMBER: 105/10/258,144

CURRENT PLILING DATE: 2002-10-17

PRIOR FILLING DATE: 2000-04-17
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PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
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Publication No. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITILE OF INVENTION: JAVELINIZATION OF PROTEINS
TITLE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: 60/197,462
FRIOR APPLICATION NUMBER: 60/197,462
PRIOR APPLICATION NUMBER: 60/197,462
PRIOR FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER: OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
                  FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: US/10/276,
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 489
LENGTHRE: FastSEQ for Windows Version 3.0
SEQ ID NO 489
LENGTH: 26
TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 493
LENGTH: 26
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ORGANISM: Homo sapiens derivative
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  TITLE OF INVENTION:
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Fublication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Phil

APPLICANT: Houghton, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

APPLICANT: New Meritons Meritons and Compositions For Heat shock

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA

FILE REFERENCE: 11746/46401

CURRENT FILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLICATION NUMBER: 60/197,642

PRIOR APPLING DATE: 2000-04-17

SOFTWARE: FESTERG for Windows Version 3.0

SEC ID NO 483

LENGTH: 26
                     APPLICANT: Al-Awgati, Gais
APPLICANT: Markw, Mark
APPLICANT: Maybew, Mark
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2002-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
SEQ ID NO 479
LENGTH: 26
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: How, Merk
APPLICANT: How, Merk
APPLICANT: How, Merk
APPLICANT: How, Merk
APPLICANT: How, Merk
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
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                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Homo sapiens derivative
US-10-258-144-479
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ORGANISM: Homo sapiens derivative
Livingston, Phil
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
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100.0%; Score 64; DB 4; Length 27; 100.0%; Pred. No. 0.09;
                                                                                        0; Indels
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Publication No. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Nothman, James
APPLICANT: Nothman, James
APPLICANT: Hoe, Mee
ITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT PILING DATE: 2002-10-17
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR PELLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 24
LENGTH: 30
                                                                                                                                                                                                                                                                       MS-10-258-147-18
Sequence 18, Application US/10258147
Sequence 18, Application US/10258147
Sequence 18, Application US/10258147
Semigral INFORMATION
APPLICANT: Rothman, James
APPLICANT: Mayhew, Mark
APPLICANT: Mos, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
PRIOR APPLICATION NUMBER: 60/197,462
PRIOR PILING DATE: 2000-10.7
NUMBER OF SEQ. ID NOS: 47
SOFTWARE: FRASEQ for Windows Version 3.0
                                                                                        0; Mismatches
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ORGANISM: Aritifical Sequence
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                   Query Match
Best Local Similarity 100...
8; Conservative
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Best Local Similarity 100...
Loc 8; Conservative
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US-10-258-144-463
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LENGTH: 30
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US-10-258-144-463
is Sequence 463, Application US/10258144
is bublication No. US20040101532A1
is GENERAL INPORMATION:
is GENERAL INPORMATION:
is APPLICANT: Houghton, Alan
is APPLICANT: Al-Awgati, Qais
is APPLICANT: MayNew, Mark
is APPLICANT: MayNew, Mark
is APPLICANT: MayNew, Mark
is APPLICANT: MayNew, Mark
is APPLICANT: Hos, Mee
it TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
is FILE REFERENCE: 11746/4401
is CURRENT APPLICATION NUMBER: US/10/258,144
is CURRENT APPLICATION NUMBER: 60/197,642
is PRIOR FILING DATE: 2000-04-17
                                                                                           FEATURE:

OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin US-10-258-147-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin
US-10-258-147-20
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100.0%; Score 64; DB 4; Length 27;
Best Local Similarity 100.0%; Pred. No. 0.09;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                   Query Match 100.0%; Score 64; DB 4; Length 27; Best Local Similarity 100.0%; Pred. No. 0.09; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20, Application US/10258147
; Publication No. US20040043419A1
; Publication No. US20040043419A1
; GENERAL INFORMATION:
; APPLICANT: Mayhew, Mark
; APPLICANT: Mos, Mark
; APPLICANT: Hoe, Mark
; TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
; CURRENT APPLICATION NUMBER: US/10/258,147
; CURRENT FILING DATE: 2002-10-17
; PRIOR FILING DATE: 2002-10-17
; RICH REPLICATION NUMBER: 60/197,462
; FRIOR FILING DATE: 200-04-17
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 20
; LENGTH: 27

TAVED: DENGTH: 27
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 463
LENGTH: 27
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                                            TYPE: PRT ORGANISM: Aritifical Sequence
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, OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin US-10-258-147-18
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; OTHER INFORMATION: javelinized peptide of Herpes Simplex virus
US-10-258-147-24
                                                           100.0%; Score 64; DB 4; Length 30; 100.0%; Pred. No. 0.098; rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 64; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.098;
Matches 8; Conservative 0; Mismatches 0; Indels
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US-10-258-144-213
; Sequence 213, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
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ORGANISM: Aritifical Sequence
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  FEATURE:
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APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: How, Nee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR PRILING DATE: 2000-04-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: FastSEQ for Windows Version 3.0
SEQ ID NO 276
LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Al-Awgati, Dais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: MayNew, Mark
APPLICANT: MayNew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746,4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 06/197,642
PRIOR APPLICATION NUMBER: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 281
LENGTH: 30
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NAME/KEY: VARIANT
LOCATION: (20)...(22)
OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
                                                                                                                            Sequence 276, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 281, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION: APPLICANT: Houghton, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens derivative
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1 HWDFAWPW 8
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HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF ILLING DATE: 2002-10-17
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 213
LENGTH: 31
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US-10-258-147-27
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Query Match
100.0%; Score 64; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.098;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                             Sequence 27, Application US/10258147
Fublication No. US20040043419A1
Fublication No. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
TILLE OF INVENTION NUMBER: US/10/258,147
CURRENT PILING DATE: 2002-10-17
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR PILING DATE: 2000-04-17
FRIOR APPLICATION NUMBER: 60/197,462
FRIOR FILING DATE: 2000-04-17
SROFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 27
LENGTH: 31
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ORGANISM: Homo sapiens derivative FEATURE:
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; Sequence 216. Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION;
APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
; APPLICANT: Al-Awqati, Qais
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APPLICANT: Al-Awqati, Uais
APPLICANT: Al-Awqati, Uais
APPLICANT: Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati, Al-Awqati
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: MATHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA
FILE REFERENCE: 11746/464610
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 221
LENGTH: 31
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           Score 64; DB 4; Length 31; Pred. No. 0.1; 0; Mismatches 0; Indels
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
GTHER INFORMATION: Any smino acid residue FEATURE:
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CCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-216
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                   100.0%;
           Query Match
Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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PERATURE:

NAME/KET: VARIANT

LOCATION: (21)...(11)

NAME/KET: VARIANT

LOCATION: (21)...(13)

STATURE:

NAME/KET: VARIANT

LOCATION: (21)...(13)

STATURE:

NAME/KET: VARIANT

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Sequence 241, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Phil

APPLICANT: Al-Awqati, Qais

APPLICANT: How, Mark

APPLICANT: How, Mark

APPLICANT: How, Mark

TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA

TITLE OF INVENTION: PROTEIN WEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT FILING DATE: 2000-04-17

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 241

LENGTH: 31

TWORE. DEPT.
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APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: How, Mark
APPLICANT: How, Mer
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA,
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
FEATURE:
NAME/KEY: VARIANT
LOCATION: (2)...(23)
LOCATION: (21)...(23)
COTHER INFORMATION: Any amino acid residue
US-10-258-144-241
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LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 246
LENGTH: 31
TYPE: PRT
ORGANISM: Homo sapiens derivative
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Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
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OTHER INFORMATION: Any amino acid
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Best Local Similarity 100.0
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TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NOS: 31
LENGTH: 31
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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NAME/KEY: VARIANT
LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 236, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.0%;
Matches 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
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1 HWDFAWPW
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FEATURE:
NOAME/KEY: VARIANT
NOAME/TON: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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COCATION: (21)...(23)
COTHER INFORMATION: Any amino acid residue
US-10-259-144-256
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; LOCATION: (21)...(23)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-261
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OTHER INFORMATION: Any amino acid residue
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FURDICARTION NO. US20040101532A1

APPLICANT: HOUGHTON, Alan

APPLICANT: Al-Angati, Oals

APPLICANT: Al-Angati, Oals

APPLICANT: Mayhew, Mark
                  TYPE: PRT ORGANISM: Homo sapiens derivative
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ORGANISM: Homo sapiens derivative
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NAME/KEY: VARIANT
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is Sequence 251, Application US/10258144

is Publication No. US20040101532A1

is GENERAL INFORMATION:

A PPLICANT: Houghton, Alan

A APPLICANT: Livingston, Phil

A APPLICANT: Al-Awgati, Qais

A APPLICANT: Hoe, We

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

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ITILE OF INVENTION: METHODS AND COMPOSITIONS

CURRENT PILING DATE: 2002-01-17

PRIOR APPLICATION NUMBER: 2002-01-17

PRIOR APPLICATION NUMBER: 2000-04-17

NUMBER: OF SEQ ID NOS: 502

SOFTWARE: PASESED for Windows Version 3.0

SEQ ID NO 251

LENGTH: 31

"WORLD NO 251

LENGTH: 31
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; Sequence 256, Application US/10258144
; bublication No. US20040101532A1
; GENERAL INFORMATION;
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
; FILE REPERENCE: 11746/4401
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT FILING DATE: 2002-10-17
; PRIOR FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: FastSEQ for Windows Version 3.0
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                                     Query Match 100.0%; Score 64; DB 4; Length 31; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
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NAME/KEY: VARIANT
LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-251
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OTHER INFORMATION: Any amino acid residue
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                   1 HWDFAWPW 8
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1 HWDFAWPW 8
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NAME/KEY: VARIANT
US-10-258-144-246
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US-10-258-144-261

i Sequence 261, Application US/10258144

j Publ.cation No. US20040101532A1

j GENERAL INFORMATION:

i APPLICANT: Houghton, Alan

j APPLICANT: Livingeton, Phil

APPLICANT: M-Awqati, Qais

j APPLICANT: Mayhew, Mark

i APPLICANT: Hoe, We

j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

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j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

j TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

j FILE REPERENCE: 11146/4640

j FILE REPERENCE: 11146/4640

j PRIOR FILING DATE: 2000-04-17

j PRIOR FILING DATE: 2000-04-17

j PRIOR FILING DATE: 2000-04-17

j NUMBER: FastSEQ for Windows Version 3.0

j SEQ ID NO 261

LENGTH: HOR FILE OF INVENTION WINDS AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND COMPOSITION OF THE METHOD AND
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100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
Query Match
100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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1 HWDFAWPW 8
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1 HWDFAWPW 8
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APPLICANT: HOUGHLOD, Alan

APPLICANT: HOUGHLOD, Phil

APPLICANT: Al-awqari, Qais

APPLICANT: Al-awqari, Qais

APPLICANT: Al-awqari, Qais

APPLICANT: Al-awqari, Qais

APPLICANT: Mayhew, Mark

APPLICANT: Hoe, Mee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/4401

CURRENT APPLICATION NUMBER: 00/197,642

PRIOR PLING DATE: 2000-04-17

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTHARE: FESTESEQ for Windows Version 3.0

SEQ ID NO 291

LENGTH: 31
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APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 286
LENGTH: 31
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100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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NAME/KEY: VARIANT

LOCATION: (21)...(23)

W. CTHER INFORMATION: Any amino acid residue
US-10-258-144-286
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NAME/KEX: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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COTHER INFORMATION: Any amino acid residue
US-10-258-144-291
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; Sequence 291, Application US/10258144
; Publication No. US20040101532A1
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ORGANISM: Homo sapiens derivative
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Sequence 326. Application US/1028144
; Sequence 326. Application US/1028144
; Publication No. US20040101532A1
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Livingston, Alan
; APPLICANT: Al-Awqati, Qais
; APPLICANT: Mayhaw, Mark
; APPLICANT: Mayhaw, Mark
; APPLICANT: Hoe, Mee
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA; FILE REFERENCE: 11746/4401
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT APPLICATION NUMBER: 60/197,642
; PRIOR APPLICATION NUMBER: 60/197,642
; RIOR APPLICATION UNDER: 60/197,642
; RIOR APPLICATION UNDER: 500-417
; NUMBER OF SEQ ID NOS: 502
; SEQ ID NO 326
; SEQ ID NO 326
; LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100.0%; Score 64; DB 4; Length 31; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                      NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                               NAME/KEY: VARIANT
| LOCATION: (21)...(23)
| OTHER INFORMATION: Any amino acid residue
US-10-258-144-321
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OTHER INFORMATION: Any amino acid residue
FEATURE:
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LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 321 LENGTH: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Homo sapiens derivative PEATURE:
                                                                                                        ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity luv...
8; Conservative
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                                                                                   TYPE: PRT
                                                                                                                                      FEATURE:
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US-10-258-144-316

Sequence 316, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Al-Awgati, Qais
APPLICANT: How, Mark
APPLICANT: How, Mark
APPLICANT: How, Mark
APPLICANT: How, Mark
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APPLICANT: How, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: MWBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 2002-10-17
PRIOR PILING DATE: 2000-04-17
NUMBER: OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 316
LENGTH: 31
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: BROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILING DATE: 2002-10-17
PRIOR PILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
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Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                           Query Match 100.0%; Score 64; DB 4; Length 31; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
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ORGANISM: Homo sapiens derivative
FEATURE:
NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
PEATURE:
NAME/KEY: VARIANT
     i LOCATION: (21)...(23)
i OTHER INFORMATION: Any amino acid residue
US-10-258-144-301
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OTHER INFORMATION: Any amino acid residue
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1 HWDFAWPW 8
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Sequence 391, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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                                                                                                                                US-10-258-144-391
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Houghton, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 2002-10-17
PRIOR APPLICATION NUMBER: 6/197,642
PRIOR APPLICATION NUMBER: 6/197,642
NUMBER OF SEC ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 386
LENGTH: 31
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
SEQ ID NO 331
LENGTH: 31
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100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: VARIANT
LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-331
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
FEATURE:
NAME/KEY: VARIANT
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i OTHER INFORMATION: Any amino acid residue
US-10-258-144-386
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Homo sapiens derivative
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Gaps
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Amagari, Qais
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF ILVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF ILLNG DATE: 2002-10-17
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOUTHARE: FastSEQ for Windows Version 3.0
SEQ ID NO 416
LENGTH: 31
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100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches A. T. J.
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OTHER INFORMATION: Any amino acid residue
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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; OTHER INFORMATION: Any amino acid residue
US-10-258-144-391
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; Sequence 416, Application US/10258144

; Publication No. US20040101532A1

; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Homo sapiens derivative
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ORGANISM: Homo sapiens derivative
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PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 426
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Matches 8; Conserv
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                 LENGTH: 31
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; Sequence 421, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION;
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; PRICANT: Mayhew, Mark
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; PRICANT: Mayhew, Mark
; PRICANT: Mayhew, Mark
; PRICANT: MABER: US/10/258,144
; CURRENT FILING DATE: 2002-10-17
; PRIOR PILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: PESESEQ for Windows Version 3.0
; SEQ ID NO 421
; TYPE: PRICANT: MARK
; PRICANT: MARK
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US-10-258-144-426
; Sequence 426, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Livingston, Phil
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Al-Awgati, Qais
; APPLICANT: Hoe, Me
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; FILLS REPRENCE: 11746/46401
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT FILLING DATE: 2002-10-17
; PRIOR APPLICATION NUMBER: 60/197,642
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                                                                                                                                                                                                                    100.0%; Score 64; DB 4; Length 31; 100.0%; Pred. No. 0.1; Ative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens derivative FEATURE: VARIANT LOCATION: (9)...(11) OTHER INFORMATION: Any amino acid residue
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CCATION: (21)...(23)

OTHER INFORMATION: Any amino acid residue

VS-10-258-144-421
                                                                      LOCATION: (21)...(23)
OTHER INFORMATION: Any amino acid residue
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                 1 HWDFAWPW 8
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   FEATURE:
NAME/KEY: VARIANT
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US-10-258-144-431

i Sequence 431. Application US/10258144

i Publication No. US20040101532A1

i GENERAL INFORMATION:
    APPLICANT: Houghton, Alan
    APPLICANT: Livingston, Phil
    APPLICANT: Al-Awqati, Qais
    APPLICANT: Hos, Mark
    APPLICANT: Hos, Mark
    APPLICANT: Hos, Mer
    TILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
    TILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
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    TILLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
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100.0%; Pred. No. 0.1;
rative 0; Mismatches 0; Indels
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                                                                                    FEATURE:
NAME/KEY: VARIANT
CACATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
FEATURE:
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                       LOCATION: (21)...(23)
CTHER INFORMATION: Any amino acid residue US-10-258-144-426
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// OTHER INFORMATION: Any amino acid residue
US-10-258-144-431
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US-10-258-144-436
S. Sequence 436, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens derivative
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: 2002-10-17
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
SEQ ID NO 471
LENGTH: 31
                                                                        PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 31;
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100.0%; Score 64; DB 4;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0;
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OTHER INFORMATION: Any amino acid residue
FEATURE:
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; LOCATION: (21)...(23)
; OTHER INFORMATION: Any amino acid residue
US-10-258-144-466
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Publication No. US20040101532A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens derivative
FEATURE:
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ORGANISM: Homo sapiens derivative
FEATURE:
    1 HWDFAWPW
                                        1 HWDFAWPW
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                                    APPLICANT: "GOVER, FIGER
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT PILING DATE: 2001-01-7
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FRESESQ for Windows Version 3.0
SEQ ID NO 436
LENGTH: 31
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746646101
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTHARE: FastSEQ for Windows Version 3.0
SEQ ID NO 441
LENGTH: 31
TYPE: PRT
ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 64; DB 4; Length 31; 100.0%; Pred. No. 0.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   i LOCATION: (21)...(23)
i OTHER INFORMATION: Any amino acid residue
US-10-258-144-436
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i OTHER INFORMATION: Any amino acid residue
US-10-258-144-441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/10258144 o. US20040101532A1
                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens derivative
Livingston, Phil
Al-Awgati, Qais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                     Al-Awgati, Qa
Mayhew, Mark
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NAME/KEY: VARIANT
LOCATION: (9)...(
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NAME/KEY: VARIANT
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Gaps

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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 501
LENGTH: 31
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; Sequence 266, Application US/10258144
; Publication No. US20040101532A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: Hybrid antigen US-10-820-067A-919
                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.
Matches 8; Conservative
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APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hos, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN SOU2-10-17
CURRENT APPLICATION NUMBER: 60/197, 642
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 496
LENGTH: 31
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100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                             Query Match
100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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CTHER INFORMATION: Any amino acid residue
US-10-258-144-496
                                                                        NAME/KEY: VARIANT

CCATION: (21)...(23)

OTHER INFORMATION: Any amino acid residue
US-10-258-144-471
       LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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Publication No. US20040101532A1
GENERAL INFORMATION:
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1 HWDFAWPW 8
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US-10-258-144-501
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                                                                                                                                                                                                                                                    Query Match
100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VESU-14-09
VESU-16-820-067A-919
VESU-16-820-067A-919
Sequence 919, Application US/10820067A
PUblication No. US20050214312A1
GENERAL INFORMATION:
APPLICANT: Fletchner, J.
APPLICANT: Mehte, S.
APPLICANT: Andielic, S.
APPLICANT: Andielic, S.
APPLICANT: Barber, B.
TITLE OF INVENTION: VACCINES AND IMMUNOTHERAPIES
FILE REPERBYCE: 8449-406-999
CURRENT FILING DATE: 2004-04-08
FRIOR APPLICATION NUMBER: 60/462,469
PRIOR PILING DATE: 2003-04-11
PRIOR PLING DATE: 2003-04-11
PRIOR PLLING DATE: 2003-04-11
PRIOR FLLING DATE: 2003-04-11
PRIOR FLLING DATE: 2003-04-11
PRIOR FLLING DATE: 2003-04-11
SPRIOR FLLING DATE: 2003-04-11
PRIOR FLLING DATE: 2003-04-11
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SPRIOR FLLING DATE: 2003-04-11
SPRIOR FLLING DATE: 2003-04-11
SPRIOR FLLING DATE: 2003-04-11
SPRIOR FLLING DATE: 2003-04-11
SPRIOR FLLING DATE: 2003-09-16
NUMBER OF SEQ ID NOS: 926
SOFTWARE: FRRESEQ FOR WINGOWS VERSION 4.0
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                        NAME/KEY: VARIANT
CCATION: (21)...(23)
CTHER INFORMATION: Any amino acid residue
US-10-258-144-501
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Gaps
Gaps
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APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqari, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT APPLICATION NUMBER: 2002-10-17
PRIOR PELING DATE: 2000-04-17
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PSECSEQ for Windows Version 3.0
SEQ ID NO 306
LENGTH: 32
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Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Houghton, Alan

APPLICANT: Al-Awqeti, Qais

APPLICANT: Al-Awqeti, Qais

APPLICANT: Hoe, Mee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT PILING DATE: 2002-10-17

PRIOR PPLICATION NUMBER: 60/197,642

PRIOR FILING DATE: 2000-04-17
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Mismatches
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; OTHER INFORMATION: Any amino acid residue US-10-258-144-306
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OTHER INFORMATION: Any amino acid residue
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SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 311
LENGTH: 32
TYPE: PRT
OXGANISM: Homo sapiens derivative
FEATURE:
                                                                                                                                                                                                US-10-258-144-306
... Sequence 306, Application US/10258144
... Publication No. US20040101532A1
... GENERAL INFORMATION:
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Best Local Similarity 100.0
....heg 8; Conservative
8; Conservative
                                                 1 HWDFAWPW 8
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1 HWDFAWPW 8
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NAME/KEY: VARIANT
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  Matches
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                  APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 266
LENGTH: 32
LENGTH: 32
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Publication No. US20040101532A1
Publication No. US20040101532A1
Publication No. US20040101532A1
APPLICANT: Houghton, Alan
APPLICANT: Al-Awqati, Qais
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTIEN MEDIATED IMMUNOTHERAPY OF MELANOMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 64; DB 4; Length 32;
Pred. No. 0.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DB 4; Length 32; 100.0%; Pred. No. 0.1; tive 0; Mismatches 0; Indels
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COTHER INFORMATION: Any amino acid residue
US-10-258-144-271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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Best Local Similarity 100.
Matches 8; Conservative
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Best Local Similarity
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NAME/KEY: VARIANT
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LENGTH: 32
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1 HWDFAWPW 8
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NAME/KEY: VARIANT
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US-10-258-144-351
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Sequence 336, Application US/10258144

Publication No. U320040101532A1

GENERAL INFORMATION:
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: How, Mark
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
FILE REFRERENCE
FILE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REPRENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2000-10-17
PRIOR PRILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 336
LENGTH: 33
TYPE: PRI
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Publication No. US20040101532A1
Publication No. US20040101532A1
Publication No. US20040101532A1
APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Hoe, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
                                                                                                                                                                                    Query Match 100.0%; Score 64; DB 4; Length 32; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                           FEATURE:
NAME/KEY: VARIANT
LOCATION: (12)...(24)
OTHER INFORMATION: Any amino acid residue
NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: VARIANT
LOCATION: (22)...(24)
CTHER INFORMATION: Any amino acid residue
US-10-258-144-336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 11746/46401
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HWDFAWPW 8
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                             US-10-258-144-311
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CURRENT APPLICATION NUMBER: 10/10/256,144

CURRENT APPLICATION NUMBER: 10/10/256,144

PRIOR PLING DATE: 2002-10-17

PRIOR PLING DATE: 2002-04-17

PRIOR PLING DATE: 2002-04-17

PRIOR PLING DATE: 2002-04-17

PRIOR PLING DATE: 2002-04-17

SCRANISH: Nono sapiens derivative fraging and translation of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said of the said
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Query Match 100.0%; Score 64; DB 4; Length 32; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
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OTHER INFORMATION: Any amino acid residue
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CTHER INFORMATION: Any amino acid residue
US-10-258-144-361
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; Sequence 361, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
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NAME/KEY: VARIANT
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
APPLICANT: Meyhew, Mark
APPLICANT: Meyhew, Mark
APPLICANT: Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REPERENCE: 11746/46101
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PELING DATE: 2000-64-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PESESSEQ for Windows Version 3.0
SEQ ID NO 356
LENGTH: 32
                                                                   APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Amgati, Qais
APPLICANT: Al-Amgati, Qais
APPLICANT: Al-Amgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: DATE: 2002-10-17
CURRENT APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NOS: 502
SEQ ID NO 351
LENGTH: 32
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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; NAME/KEY: VARIANT
; LOCATION: (22)...(24)
D CTHER INFORMATION: Any amino acid residue
US-10-258-144-356
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CCATION: (22)...(24)

CTHER INFORMATION: Any amino acid residue

US-10-258-144-351
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  Sequence 351, Application US/10258144
Publication No. US20040101532A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.
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Gaps
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APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Moyhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PELING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PREESEQ for Windows Version 3.0
SEQ ID NO 361
LENGTH: 32
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APPLICANT: Houghton, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Me
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: MOUBBER: 1010/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLICING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500-
SOFTWARE: FREESEQ FOR WINDOWS Version 3.0
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100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNO?
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SEQ ID NO 376
LENGTH: 32
TYPE: PRI
TYPE: PRI
TYPE: PRI
TYPE: PRI
FEATURE:
FEATURE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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CCATION: (22)...(24)

OTHER INFORMATION: Any amino acid residue

US-10-258-144-376
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1 LOCATION: (22)...(24)
2 OTHER INFORMATION: Any amino acid residue
US-10-258-144-381
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OTHER INFORMATION: Any amino acid residue
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1 HWDFAWPW 8
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Publication No. US20040101532A1

GENERAL INFORMATION;

APPLICANT: Houghton, Alan

APPLICANT: Houghton, Phil

APPLICANT: How, Mark

APPLICANT: Mayhew, Mark

APPLICANT: Al-Awqati, Qais

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2000-04-17

PRIOR PILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FEBLESC for Windows Version 3.0

SEQ ID NO 371

LENGTH: 32

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APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
IIILE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
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                                                                                                             LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
FEATURE:
NAME/KEY: VARIANT
LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-366
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORWATION: Any amino acid residue
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NAME/KEY: VARIANT
LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
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US-10-258-144-376
; Sequence 376, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
          ORGANISM: Homo sapiens derivative FEATURE: NAME/KEY: VARIANT
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US-10-258-144-381

Sequence 381, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION.

APPLICANT: Houghton, Alan

APPLICANT: Al-Awgari, Qais

APPLICANT: Al-Awgari, Qais

APPLICANT: Hoe, Mee

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: PROTHEN MEDIATED INMUNOTHERAPY OF MELANOWA

FILE REFERENCE: 11746/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17

PRIOR APPLICATION NUMBER: 60/197,642

MUMBER OF SEQ ID NOS: 502

SOTWARRE: FRACESEQ for Windows Version 3.0

SEQ ID NO 381

LENGTH: ALCOR PILING DATE: ALCORDING NUMBER: US/10/17

PRIOR APPLICATION NUMBER: 2000-04-17

MUMBER OF SEQ ID NOS: 502

SEQ ID NO 381

LENGTH: ALCORDING NUMBER: ALCORDING NUMBER: US/10/17

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Query Match
100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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0; Mismatches
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OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 411
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. Sequence 406. Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
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Publication No. US20040101532A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens derivative
                   Query Match
Best Local Similarity 100.
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Matches 8; Conservative
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1 HWDFAWPW
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NAME/KEY: VARIANT
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US-10-258-144-401
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; bublication No. UG20040101532A1
; bublication No. UG20040101532A1
; GENERAL INFORMATION;
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: MayNew, Mark
; APPLICANT: MayNew, Mark
; APPLICANT: MayNew, Mark
; APPLICANT: Al-Awqati, Qais
; APPLICANT: Hoe, Wee
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
; TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA; FILE REFERENCE: 11746/46401
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT APPLICATION NUMBER: 60/197,642
; RIOR PAPLING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SEQ ID NO 401
; LENGTH: 32
                                                                                                       APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 105/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 500
SOFTWARE: FastSEQ for Windows Version 3.0
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NAME/KEY: VARIANT
LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-396
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OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
                                         Sequence 396, Application US/10258144 Publication No. US20040101532A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens derivative
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NAME/KEY: VARIANT
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LENGTH: 32
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APPLICANT: Houghton, Phil
APPLICANT: Livingston, Phil
APPLICANT: Livingston, Phil
APPLICANT: Al-Awagati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: ROOTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PELLING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 406
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APPLICANT: Al-Awgati, Oais
APPLICANT: Al-Awgati, Oais
APPLICANT: Mayhew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
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100.0%; Score 64; DB 4; Length 32; 100.0%; Pred. No. 0.1;
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                                                                                                                                         Indels
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Gaps
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Sequence 456, Application US/20258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Livingston, Alan

APPLICANT: Livingston, Phil

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

APPLICANT: Al-Awgati, Qais

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

CURRENT APPLICANTON NUMBER: US/10/258,144

CURRENT FILING DATE: 2002-10-17

PRIOR PRILING DATE: 2000-44-17

NUMBER OF SEQ ID NOS: 502

SEQ ID NOS: 502

SEQ ID NOS: 502

SEC IT NO 456
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APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OP INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: US/10/258,144
PRIOR APPLICATION NUMBER: 0S/197,642
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PARESEQ for Windows Version 3.0
SEQ ID NO 451
LENGTH: 32
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NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
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CTHER INFORMATION: Any amino acid residue US-10-258-144-451
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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Best Local Similarity 100.0
Matches 8; Conservative
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Matches 8; Conservative
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US-10-258-144-456
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LENGTH: 32
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US-10-258-144-446

JUS-10-258-144-446

Sequence 446, Application US/10258144

Publication No. US20040101532A1

GENERAL INFORMATION:

APPLICANT: Livingeron, Alan

APPLICANT: Livingeron, Phil

APPLICANT: Mayhew, Mark

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK

CURRENT APPLICANT: US 11/46/46401

CURRENT APPLICATION NUMBER: US/10/258,144

CURRENT APPLICATION NUMBER: 60/197,642

PRIOR FILING DATE: 2000-04-17

NUMBER OF SEQ ID NOS: 502

SOFTWARE: FRASESEQ for Windows Version 3.0

SEQ ID NO 446

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100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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NAME/KEY: VARIANT
LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-411
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NAME/KEY: VARIANT

LOCATION: (22)...(24)

OTHER INFORMATION: Any amino acid residue
US-10-258-144-446
                                                                                                          FEATURE:
NAME/KEX: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
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US-10-258-144-451
; Sequence 451, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
    APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Al-Awqati, Qais
; APPLICANT: Mayhew, Mark
                                            TYPE: PRT
ORGANISM: Homo sapiens derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens derivative
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1 HWDFAWPW 8
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i LOCATION: (22)...(24)
i OTHER INFORMATION: Any amino acid residue
US-10-258-144-476
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NAME/KEY: VARIANT
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APPLICANT: LIVINGSTON, PAIL
APPLICANT: Al-Awqati, Qais
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED INMUNOTHERAPY OF MELANOMA
TITLE OF INVENTION: DESCRIPTION OF MELANOMA
FILE REFERENCE: 11746/461 US/10/258,144
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLILNG DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PSSEC for Windows Version 3.0
SEQ ID NO 476
LENGTH: 32
                                                                                                                                                                 APPLICANT: Houghton, Alan
APPLICANT: Livingston, Phil
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Al-Awgati, Qais
APPLICANT: Hoi- Meynew, Mark
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
CURRENT PELICANTON NUMBER: 105/10/258,144
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR PLILING DATE: 2000-04-17
NUMBER OF SEC ID NOS: 500-04-17
NUMBER OF SEC ID NOS: 500-04-17
NUMBER OF SEC ID NOS: 500-04-17
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i OTHER INFORMATION: Any amino acid residue
US-10-258-144-461
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OTHER INFORMATION: Any amino acid residue
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OTHER INFORMATION: Any amino acid residue
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Sequence 476, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
APPLICANT: Houghton, Alan
                                                                                              Application US/10258144
O. US20040101532A1
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ORGANISM: Homo sapiens derivative
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ORGANISM: Homo sapiens derivative
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Best Local Similarity 100.
Matches 8; Conservative
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HWDFAWPW 8
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NAME/KEY: VARIANT
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LENGTH: 32
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                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Houghton, Alan
APPLICANT: Houghton, Phil
APPLICANT: Al-Awgari, Qais
APPLICANT: Al-Awgari, Qais
APPLICANT: Al-Awgari, Qais
APPLICANT: Al-Awgari, Qais
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOWA
FILE REFERENCE: 11746/46401
CURRENT APPLICATION NUMBER: 00/10/17
PRIOR APPLICATION NUMBER: 60/197,642
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
SOFTWARE: PastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 430
US-10-258-144-486
iSequence 486, Application US/10258144
j Publication No. US20040101532A1
iGENERAL INFORMATION:
APPLICANT: Houghton, Alan
j APPLICANT: Livingston, Phil
APPLICANT: Mayhaw, Mark
APPLICANT: MAYHAW, Mark
TITLE OF INVENTION: PROTEIN METHODS AND COMPOSITIONS FOR HEAT SHOCK
TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
FILE REFERENCE: 11746/4401
CURRENT APPLICATION NUMBER: US/10/258,144
CURRENT APPLICATION NUMBER: 00/197,642
PRIOR FILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 502
                                                           ö
Query Match
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 64; DB 4; Length 32; Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: VARIANT
LOCATION: (22)...(24)
OTHER INFORMATION: Any amino acid residue
US-10-258-144-481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                                                                                                                                                    Sequence 481, Application US/10258144
Publication No. US20040101532A1
GENERAL INFORMATION:
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Gaps

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; FEATURE:
; OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin
US-10-258-147-21
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100.0%; Score 64; DB 4; Length 38;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 8; Conservative 0; Mismatches 0; Indels
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,462
PRIOR PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PASTSEQ for Windows Version 3.0
SEQ ID NO 21
                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Aritifical Sequence
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LENGTH: 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 491, Application US/10258144
; Sequence 491, Application US/10258144
; Publication No. US20040101532A1
; GENERAL INFORMATION:
; APPLICANT: Houghton, Alan
; APPLICANT: Livingston, Phil
; APPLICANT: Livingston, Phil
; APPLICANT: Mayhew, Mark
; APPLICANT: Mayhew, Mark
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; TITLE OF INVENTION: PROTEIN MEDIATED IMMUNOTHERAPY OF MELANOMA
; FILE REFERENCE: 11746/46401
; CURRENT APPLICATION NUMBER: US/10/258,144
; CURRENT PILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 502
; SOFTWARE: FBSLSEQ for Windows Version 3.0
; SEQ ID NO 491
; LENGTH: 32
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100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                        FEATURE:
NAME/KEY: VARIANT
LOCATION: (9)...(11)
OTHER INFORMATION: Any amino acid residue
                                                                                                                                                                                              FEATURE:
NAME/KEY: VARIANT
LOCATION: (22)...(24)
US-10-258-144-486
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (22)...(24)
COTHER INFORMATION: Any amino acid residue
US-10-258-144-491
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                               TYPE: PRT
ORGANISM: Homo sapiens derivative
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1 HWDFAWPW 8
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                    SEQ ID NO 486
LENGTH: 32
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100.0%; Score 64; DB 4; Length 100;
Best Local Similarity 100.0%; Pred. No. 0.26;
Matches 8; Conservative 0; Mismatches 0; Indels
Sequence 28, Application US/10258147
PUDLICALION NO. US2004004319A1
GENERAL INFORMATION;
JAPPLICANT: Rothman, James
JAPPLICANT: Rothman, James
JAPLICANT: Hoe, Mark
APPLICANT: Hoe, Mark
GURRENT POST 1746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/197,462
NUMBER OF SEG ID NOS: 47
SOFTWARE: FRASEQ for Windows Version 3.0
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Publication No. US20040043419A1

GENERAL INFORMATION:
APPLICANT: Rothman, James
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147

CURRENT APPLICATION NUMBER: 002-10-17

PRIOR APPLICATION NUMBER: 60/197,462

PRIOR FILING DATE: 2000-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Aritifical Sequence
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; OTHER INFORMATION: javelinized peptide of Gallus gallus ovalbumin US-10-258-147-31
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 108
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ORGANISM: Aritifical Sequence
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                                                                                                                                               1 HWDFAWPW 8
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1 HWDFAWPW 8
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100.0%; Score 64; DB 4; Length 100;
Best Local Similarity 100.0%; Pred. No. 0.26;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 64; DB 4; Length 103;
Best Local Similarity 100.0%; Pred. No. 0.26;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                              RESULT 435
US-10-258-147-29
iS-10-258-147-29
sequence 29, Application US/10258147
publication No. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Mayhew, Mark
APPLICANT: Mayhew, Mark
TILLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT FILING DATE: 2002-10-17
CURRENT FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 29
LENGTH: 103
TYDED
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APPLICANT: Rothman, James
APPLICANT: Mayhew, Mark
APPLICANT: Hoc, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REFERENCE: 11746/46276
CURRENT APPLICATION NUMBER: US/10/258,147
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 31
LENGTH: 103
    NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 30
LENGTH: 100
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Publication No. US20040043419A1
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ORGANISM: Aritifical Sequence
FEATURE:
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ORGANISM: Aritifical Sequence
                                                                                   TYPE: PRT
ORGANISM: Aritifical Sequence
                                                                                                                                                                                                                                                                                                                   93 HWDFAWPW 100
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1 HWDFAWPW
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                                                                                                                         FEATURE:
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US-10-258-147-33
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Query Match
100.0%; Score 64; DB 4; Length 103;
Best Local Similarity 100.0%; Pred. No. 0.26;
Matches 8; Conservative 0; Mismatches 0; Indels
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US-10-258-147-33

Sequence 33, Application US/10258147

Publication No. US20040043419A1

GENERAL INFORMATION

APPLICANT: Rothman, James

APPLICANT: Rothman, JaweLINIZATION OF PROTEINS

TITLE OF INVENTION: JAVELINIZATION OF PROTEINS

TITLE OF INVENTION: JAVELINIZATION OF PROTEINS

CURRENT PLILING DATE: 2002-10-17

CURRENT FILING DATE: 2002-04-17

NUMBER OF SEQ ID NOS: 47

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 33

LENGTH: 111

THENGTH: 111
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US-10-258-147-32
Sequence 32, Application US/10258147
FUBLICACION NO. US20040043419A1
GENERAL INFORMATION:
APPLICANT: Rothman, James
APPLICANT: Maybew, Mark
APPLICANT: Hoe, Mee
TITLE OF INVENTION: JAVELINIZATION OF PROTEINS
FILE REPERENCE: 11746/46276
CURRENT FILING DATE: 2000-10-17
CURRENT FILING DATE: 2000-0-17
FRIOR FILING DATE: 2000-0-17
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Tue Mar 28 10:55:31 2006

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1 HWDFAWPW 8 |||||||| 1 HWDFAWPW 8

Search completed: March 24, 2006, 12:48:25 Job time : 168 secs

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Sequence 72, Appl
Sequence 74, Appl
Sequence 74, Appl
Sequence 75, Appl
Sequence 3297, Ap
Sequence 3297, Ap
Sequence 5697, Ap
Sequence 5697, Ap
Sequence 25808, A
Sequence 25808, A
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Sequence 25808, A
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Sequence 21848, A
Sequence 933, App
Sequence 80, Appl
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10693, A
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364, App
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Sequence 31887, A
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                                                                          March 24, 2006, 08:04:28; Search time 75 Seconds (without alignments) 3.145 Million cell updates/sec
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8: /cgn2 6/ptodata/2/pubpaa/US11 NEW PUB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-10-798-579A-10
US-11-054-281-72
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US-11-054-281-74
US-11-054-281-74
US-11-054-281-75
US-11-054-281-75
US-11-091-324-3
US-11-087-099-3297
US-11-087-099-5697
US-11-096-568A-25808
US-11-096-568A-25808
US-11-096-568A-25807
US-11-096-568A-25807
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US-11-096-568A-25807
US-11-096-568A-25807
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US-11-096-568A-31887
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1-087-099-10693
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                                                                                                                                                                                                                                174681 seqs, 29487097 residues
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Maximum Match 100%
Listing first 45 summaries
                                                     OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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                                                                               Run on:
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No.
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Sequence
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US-11-096-568A-31885
US-11-096-568A-17900
US-11-096-568A-17899
US-11-096-568A-17899
US-11-096-568A-17899
US-09-978-360A-480
US-09-978-360A-681
US-11-056-558A-312
US-09-978-360A-681
US-11-052-55AA-312
US-11-096-568A-1784
US-11-218-281-23
US-11-218-281-23
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Sequence 21848, Application US/11096568A

Publication No. US20060048240A1

Publication No. US20060048240A1

GENERAL INFORMATION:

APPLICANT: Alexandrov, Nickolai et al.

TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides F; TITLE OF INVENTION: Therby

TITLE OF INVENTION: Therby

TITLE OF INVENTION: Therby

CURRENT APPLICATION NUMBER: US/11/096,568A

CURRENT APPLICATION NUMBER: US/11/096,568A

CURRENT PILING DATE: 2005-04-01

NUMBER OF SEQ ID NOS: 34471

SEQ ID NO 21848

LENGTH: 460
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Sequence 31886, A Sequence 31886, A Sequence 8, Appl 1 Sequence 17899, A Sequence 17899, A Sequence 17899, A Sequence 192, App Sequence 681, App Sequence 5352, App Sequence 5352, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 23, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 29, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App I Sequence 20, App 
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23;
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; Sequence 1920, Application US/10492570
; Publication No. US2060057666A1
; GENERAL INFORMATION:
    APPLICANT: Zhang, Jian
    TITLE OF INTENTION: A HUMAN G PROTEIN COUPLED RECEPTOR
    TILE REFERENCE: PB0180
    CURRENT APPLICATION NUMBER: US/10/492,570
    CURRENT APPLICATION NUMBER: US 60/329,000
    PRIOR APPLICATION NUMBER: US 60/329,000
    PRIOR FILING DATE: 2001-10-12
    NUMBER OF ENQ ID NOS: 1926
    SOFTWARE: Aeomica Sequence Listing Engine
    SEQ ID NO 1920
    LENGTH: 126
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ORGANISM: Rhodobacter capsulatus
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NAME/KEY: misc_feature
LOCATION: (1)..(460)
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257 DYGWPW 262
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Sequence 80. Application US/09995493

GENERAL INFORMATION:

APPLICANT: Handfield, Martin

APPLICANT: Hillman, Jeffrey

APPLICANT: Frogulake-Pox, Ann

ITILE OF INVENTION: In the Diagnosis, Treatment, and Monitoring of Periodontal Disea
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APPLICANT: Glan, Xiachong B.
APPLICANT: Wang, Zhiwei
APPLICANT: Wehrman, Tom
APPLICANT: Wehrman, Tom
APPLICANT: Zhang, Jie
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Drmanac, Radoje T.
ITLE OP INVENTION: NOVEL NOVALINGE: US/11/000,463
CURRENT APPLICATION NUMBER: US/11/000,463
CURRENT APPLICATION NUMBER: US/21,265
PRIOR PILING DATE: 2001-01-25
PRIOR PILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR PILING DATE: 2000-07-17
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-09-15
NUMBER OF SEC ID NOS: 944
SOPTWARE: PastSEQ for Windows Version 3.0
SEQ ID NOS: 944
SEQ ID NOS: 944
TUDES: DATE
                                                                 Query Match 60.9%; Score 39; DB 7; Length 460; Best Local Similarity 57.1%; Pred. No. 63; Matches 4; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 38; DB 7; Length 62;
Pred. No. 18;
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, OTHER INFORMATION: Ceres Seq. ID no. 12406509
US-11-096-568A-21848
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; Publication No. US20050266423A1
; GENERAL INFORMATION:
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Best Local Similarity 57.1%;
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Tang, Y Tom
APPLICANT: Liu, Chenghua
APPLICANT: Asundi, Vinod
APPLICANT: Chen, Rui-hong
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US-11-000-463-933
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6 WPWPWPW 12
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US-10-887-475B-10
i Sequence 10, Application US/10887475B
i Sequence 10, Application US/10887475B
i Publication No. US20060015973A1
i GENERAL INFORMATION:
i APPLICANT: SHINOZAKI, Kazuko;
i APPLICANT: SASUGA, Mie;
i APPLICANT: SASUGA, Mie;
i TITLE OF INVENTION: Environmental stress-tolerant plants
i TITLE OF INVENTION: Environmental stress-tolerant plants
i TITLE OF INVENTION NUMBER: US/10/887,475B
CURRENT PPLILNG DATE: 2004-07-09
i PRIOR APPLICATION NUMBER: US 09/301,217
i PRIOR FILING DATE: 1999-04-28
i PRIOR FILING DATE: 1999-04-28
i PRIOR FILING DATE: 1999-04-28
i PRIOR FILING DATE: 1998-10-14
i NUMBER OF SEQ ID NOS: 75
i SEQ ID NO 10
i TYPE: PRI
i ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 330;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ŝ
                                                                                                                                                                                                                                                                                                                                                                                  ) ORGANISM: Actinobacillus actinomycetemcomitans US-09-995-493-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.4%; Score 38; DB
83.3%; Pred. No. 47;
tive 0; Mismatches
FILE REFERENCE: MBHB01-662
CURRENT APPLICATION NUMBER: US/09/995,493
CURRENT FILING DATE: 2001-11-28
NUMBER OF SEQ ID NOS: 234
SOFTWARE: Patentin version 3.0
SEQ ID NO 80
LENGTH: 208
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ORGANISM: Arabidopsis thaliana
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Best Local Similarity 83.3
Matches 5; Conservative
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Gaps

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Publication No. US20060013813A1

GRERRAL INFORMATION:

APPLICANT: Mezes et al.

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same

FILE REFERENCE: 21402-240CIP

CURRENT FILING DATE: 2001-01-11

PRIOR FILING DATE: 2001-01-11

PRIOR PILING DATE: 2001-09-10

PRIOR APPLICATION NUMBER: 60/261,013

PRIOR PILING DATE: 2001-09-10

PRIOR PILING DATE: 2001-09-10

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

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PRIOR PLING DATE: 2001-01-11
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100.0%; Pred. No. 1.6e+0:
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           59.4%; bcc. No. 1.00.0%; Pred. No. 1.00.0%; Pred. No. 1.00.0%; Mismatches
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,029
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/313,170
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2002-01-11
NUMBER OF SEQ ID NOS: 324
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 14
LENGTH: 967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-11-054-281-75; Sequence 75, Application US/11054281; Sequence 75, Application WG20060013813A1; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 5; Conservative
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Matches 5; Conservative
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US-11-054-281-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          809 WDPAW 813
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Sequence 72, Application US/11054281

Fublication No. US20060013813A1

GENERAL INFORMATION:
TILE OF INVENTION: Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-240CTB

CURRENT FILING DATE: 2005-02-08

FRIOR PAPLICATION NUMBER: 60/261,014

PRIOR APPLICATION NUMBER: 60/261,018

PRIOR APPLICATION NUMBER: 60/261,018

PRIOR APPLICATION NUMBER: 60/261,018

PRIOR APPLICATION NUMBER: 60/261,018

PRIOR FILING DATE: 2001-09-10

PRIOR FILING DATE: 2001-09-10

PRIOR FILING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

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PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR APPLICATION NUMBER: 6/261,029

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR APPLICATION NUMBER: 10/044,564

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

PRIOR PLING DATE: 2001-01-11

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CURRENT APPLICATION NUMBER: 205-02-08
PRIOR PRING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: 60/261,014
PRIOR APPLICATION NUMBER: 60/261,018
PRIOR FILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,018
PRIOR PILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR FILING DATE: 2001-01-11
                                                                      Score 38; DB 6; Length 330;
Pred. No. 68;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 966;
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Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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; Sequence 14, Application US/11054281
; Publication No. US20060013813A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Orycctolagus cuniculus US-11-054-281-72
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Best Local Similarity 100.0%;
Matches 5; Conservative 0
                                      Query Match
Best Local Similarity 66.7-
                                                                                                                                                                                                                                            257 DYGWPW 262
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808 WDFAW 812
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            US-10-887-475B-10
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APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450) B EP
CURREY APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
SUD ID NOS: 12464
LENGTH: 370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 57.8%; Score 37; DB 7; 66.7%; Pred. No. 1e+02; tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                  57.8%;
       Publication No. US20060041961A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 66.,
Lac 4, Conservative
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Best Local Similarity 66.7-
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Best Local Similarity 66.7°
Matches 4; Conservative
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                                                                                                                                                                                                                                            ; TYPE: PRT
; ORGANISM: Glycine max
US-11-087-099-9967
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216 HWEFLW 221
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TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same FILE REFERENCE: 21402-240CIP
CURRENT PRILING DATE: 2005-208
PRIOR APPLICATION NUMBER: 05/261,014
PRIOR PELLING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-09-10
PRIOR PELLING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR PELLING DATE: 2001-01-11
PRIOR PELLING DATE: 2001-01-11
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PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-08-17
PRIOR PILING DATE: 2001-08-17
PRIOR PILING DATE: 2002-01-11
NUMBER OF SEQ ID NOS: 22-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3, Application US/11181234; Sequence 3, Application US/11181234; Publication No. US20060021075A1; Sequence 3, Application No. US20060021075A1; GENERAL INFORMATION:
; APPLICANT: WANG, CHYUNG-RU; TITLE OF INVENTION: GROUP 1 CD1 TRANSGENIC MICE AND THEIR; TITLE OF INVENTION: USES; FILE REFERRINGE: 2117,0001U2; CURRENT APPLICATION NUMBER: US/11/181,234; CURRENT FILING DATE: 2005-07-14; PRIOR PILING DATE: 2005-07-15; NUMBER OF SEQ ID NOS: 7; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 3; SEQ ID NO 3; LENTH: 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        59.4%; Score 38; DB 7; Length 967; 100.0%; Pred. No. 1.6e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence:/note OTHER INFORMATION: Synthetic Construct
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Best Local Similarity 42.9%; Pred. No. 79;
Matches 6; Conservative 0; Mismatches
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US-11-087-099-9967

Sequence 9967, Application US/11087099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57 WDSNSSTIVFLWPW 70
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Best Local Similarity 100.
Matches 5; Conservative
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US-11-054-281-75
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US-11-181-234-3
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Score 37; DB 7; Length 370;
Pred. No. 1e+02;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 371;
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66.7%; Pred. No. 1e+02;
cive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5697, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement;
FILE REFERENCE: 38-21(53450) B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 5697
                                                                                                                                                                                                                                 Sequence 3297, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450) B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 3297
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220 HWEFLW 225
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RESULT 15
US-11-087-099-6495
i Sequence 6495, Application US/11087099
i Publication No. US20060041961A1
i GENERAL INFORMATION:
i APPLICANT: Abad, Mark S. et al.
i TILE OF INVENTION: Genes and Uses for Plant Improvement
i TILE OF INVENTION: 38-21(53450)B EP
i CURRENT APPLICATION NUMBER: US/11/087,099
i CURRENT PILING DATE: 2005-03-22
i SEQ ID NO 6495
i LENGTH: 383
i TYPE: PRT
i ORGANISM: Glycine max
US-11-087-099-6495
                                                                                                                                                                                                                                                                                                                                                                                     Query Match 57.8%; Score 37; DB 7; Length 383; Best Local Similarity 66.7%; Pred. No. 1.18+02; Matches 4; Conservative 1; Mismatches 1; Indels
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229 HWEFLW 234
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